

OM protein - protein search, using sw model	GenCore version 5.1.4.p5_4578
Run on:	April 22, 2003, 15:27:27 ; (without alignments)
Scoring table:	OLIGO
Searcher:	Gapop 60.0 , Gapext 60.0
Word size :	0
Total number of hits satisfying chosen parameters:	908470
Minimum DB seq length:	0
Maximum DB seq length:	2000000000
Post-processing: Listing first 150 summaries	
Database :	A_Geneseq_101002:*
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6:	/SIDS2/gcdata/geneseq/geneseqp-emb1/AA1985.DAT:*
7:	/SIDS2/gcdata/geneseq/geneseqp-emb1/AA1986.DAT:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query	%	Match Length	DB ID	Description
1	1001	100.0	1001	22	AAB3533	Human TR13 receptor
2	877	87.6	1013	21	AAB26179	Human CASB619 prot
3	708	70.7	1013	22	AAB12190	Human PRO9850 poly
4	704	70.3	750	22	AAB35328	Human TR13 receptor
5	675	67.4	1013	22	AAB38345	Amino acid sequenc
6	662	66.1	911	22	AAB38351	Amino acid sequenc
7	621	62.0	870	22	AAB38351	Amino acid sequenc
8	460	45.0	495	20	AAY5972	Human endometrium
9	249	24.9	383	22	AAB38353	Amino acid sequenc
10	216	21.6	372	22	AAB85768	Human seven-transm

11	150	15.0	208	21	AAB53442	Human colon cancer
12	147	14.7	22	AAB83849	Peptide fragment o	
13	117	20.9	22	AAB83852	Amino acid sequenc	
14	96	9.6	21	AAB26180	Human CASB619 prot	
15	56	5.6	22	AAB83847	Peptide fragment o	
16	52	5.2	22	AAB83848	Peptide fragment o	
17	45	4.5	22	AAB83846	Peptide fragment o	
18	38	3.8	20	AAY2274	Human 5', EST secre	
19	21	2.1	22	AAB70256	TR16-long receptor	
20	15	1.5	22	AAB48372	Human SEC5 protein	
21	15	1.5	22	AAB48377	Human SEC10 protein	
22	15	1.5	22	AAB70255	TR16-short recepto	
23	14	1.4	22	AAB35681	Peptide #7187 enco	
24	14	1.4	22	AAB24346	Protein #6345 enco	
25	14	1.4	22	AAB70285	Human brain expres	
26	14	1.4	22	AAB7033	Human bone marrow	
27	14	1.4	22	AAM19811	Peptide #62245 enco	
28	14	1.4	22	AAM3257	Peptide #7294 enco	
29	14	1.4	23	ABG42877	Human peptide enco	
30	14	1.4	22	AAU21345	Human novel foetal	
31	14	1.4	22	AAB7028	Peptide #29. Unid	
32	10	1.0	21	AAB27114	Human CASB619 prot	
33	10	1.0	21	AAB27115	Human CASB619 prot	
34	10	1.0	21	AAB27116	Human CASB619 prot	
35	10	1.0	21	AAB27117	Human CASB619 prot	
36	10	1.0	21	AAB27118	Human CASB619 prot	
37	10	1.0	21	AAB27119	Human CASB619 prot	
38	10	1.0	21	AAB27120	Human CASB619 prot	
39	10	1.0	21	AAB27121	Human CASB619 prot	
40	10	1.0	21	AAB27122	Human CASB619 prot	
41	10	1.0	21	AAB27123	Human CASB619 prot	
42	10	1.0	21	AAB27124	Human CASB619 prot	
43	10	1.0	21	AAB27125	Human CASB619 prot	
44	10	1.0	21	AAB27126	Human CASB619 prot	
45	10	1.0	21	AAB27128	Human CASB619 prot	
46	10	1.0	21	AAB27129	Human CASB619 prot	
47	10	1.0	21	AAB27130	Human CASB619 prot	
48	10	1.0	21	AAB27131	Human CASB619 prot	
49	10	1.0	21	AAB27132	Human CASB619 prot	
50	10	1.0	21	AAB27133	Human CASB619 prot	
51	10	1.0	21	AAB27134	Human CASB619 prot	
52	10	1.0	21	AAB27135	Human CASB619 prot	
53	10	1.0	21	AAB27136	Human CASB619 prot	
54	10	1.0	21	AAB27137	Human CASB619 prot	
55	10	1.0	21	AAB27138	Human CASB619 prot	
56	10	1.0	21	AAB27139	Human CASB619 prot	
57	10	1.0	21	AAB27140	Human CASB619 prot	
58	10	1.0	21	AAB27141	Human CASB619 prot	
59	9	0.9	21	AAB26111	Human CASB619 prot	
60	9	0.9	21	AAB26182	Human CASB619 prot	
61	9	0.9	21	AAB26183	Human CASB619 prot	
62	9	0.9	21	AAB26184	Human CASB619 prot	
63	9	0.9	21	AAB26185	Human CASB619 prot	
64	9	0.9	21	AAB26186	Human CASB619 prot	
65	9	0.9	21	AAB26187	Human CASB619 prot	
66	9	0.9	21	AAB26188	Human CASB619 prot	
67	9	0.9	21	AAB26190	Human CASB619 prot	
68	9	0.9	21	AAB26191	Human CASB619 prot	
69	9	0.9	21	AAB26192	Human CASB619 prot	
70	9	0.9	21	AAB26193	Human CASB619 prot	
71	9	0.9	21	AAB26194	Human CASB619 prot	
72	9	0.9	21	AAB26195	Human CASB619 prot	
73	9	0.9	21	AAB26196	Human CASB619 prot	
74	9	0.9	21	AAB26197	Human CASB619 prot	
75	9	0.9	21	AAB26198	Human CASB619 prot	
76	9	0.9	21	AAB26200	Human CASB619 prot	
77	9	0.9	21	AAB26201	Human CASB619 prot	
78	9	0.9	21	AAB26202	Human CASB619 prot	
79	9	0.9	21	AAB27101	Human CASB619 prot	
80	9	0.9	21	AAB27102	Human CASB619 prot	
81	9	0.9	21	AAB27103	Human CASB619 prot	
82	9	0.9	21	AAB27104	Human CASB619 prot	
83	0.9	0.9	21	AAB27105	Human CASB619 prot	

84	9	9	21	AAB27106	AAB35333	AAB35333
85	9	9	9	AAB27107	AAB35333	standard; Protein; 1001 AA.
86	9	9	9	AAB27108	XX	
87	9	9	9	AAB27109	AC	AAB35333;
88	9	9	9	AAB27110	XX	
89	9	9	9	AAB27111	DT	08-MAY-2001 (first entry)
90	9	9	9	AAB27112	XX	
91	9	9	9	AAB27142	DE	Human TR13 receptor protein SEQ ID NO: 40.
92	9	9	9	AAB27149	XX	
93	9	9	9	ABB8905	KW	Human; tumour necrosis factor receptor; TR13; TR14; infection; cancer; autoimmune disease; allergy; inflammatory disease; graft rejection; apoptosis; cardiovascular disease; aneurysm.
94	9	9	9	ABB28896	KW	
95	9	9	9	AAM95554	KW	
96	9	9	9	AAM72123	KW	
97	9	9	9	AAM72123	KW	
98	9	9	9	AAM72123	KW	
99	9	9	9	AAM72123	KW	
100	9	9	9	AAM72123	KW	
101	9	9	9	AAM72123	KW	
102	8	0.9	8	AAM72123	KW	
103	8	0.8	27	AAY78753	XX	
104	8	0.8	27	AAY78753	XX	
105	8	0.8	60	ABB38686	PR	16-JUL-1999; 990US-0144087.
106	8	0.8	60	ABB38686	PR	18-AUG-1999; 990US-0149450.
107	8	0.8	60	ABB23759	PR	20-AUG-1999; 990US-0149712.
108	8	0.8	60	ABB23759	PR	10-SEP-1999; 990US-0153089.
109	8	0.8	60	AAM21249	XX	(HUMA-) HUMAN GENOME SCI INC.
110	8	0.8	60	ABG11680	PA	
111	8	0.8	64	AAM2925	XX	
112	8	0.8	64	ABG12760	PA	
113	8	0.8	74	ABG12760	XX	
114	8	0.8	74	ABG12760	XX	
115	8	0.8	22	AAM50366	DR	25-JAN-2001.
116	8	0.8	241	AAG27726	XX	
117	8	0.8	241	AAG4479	XX	
118	8	0.8	241	AAG4479	XX	
119	8	0.8	308	9 AAB8167	XX	
120	8	0.8	308	10 AABP4810	XX	
121	8	0.8	309	10 AAP35200	PT	
122	8	0.8	309	21 AAY45051	PT	
123	8	0.8	309	21 AAY45052	PT	
124	8	0.8	309	21 AAY45053	PT	
125	8	0.8	309	21 AAY45054	PT	
126	8	0.8	309	21 AAY45055	PT	
127	8	0.8	309	21 AAY45056	PT	
128	8	0.8	309	21 AAY45057	PT	
129	8	0.8	309	21 AAY45058	PT	
130	8	0.8	309	21 AAY45059	PT	
131	8	0.8	309	21 AAY45060	PT	
132	8	0.8	309	21 AAY72291	PT	
133	8	0.8	309	22 AAG79250	PS	
134	8	0.8	309	22 AAG79251	XX	
135	8	0.8	309	22 AAG79252	XX	
136	8	0.8	309	22 AAG79253	CC	The present invention provides the protein and coding sequences of the human tumour necrosis factor receptors TR13 and TR14. These sequences are useful in the diagnosis and treatment of many diseases, including cancer, autoimmune diseases, cardiovascular disorders, allergies, neurodegenerative diseases, graft rejection, inflammation, aneurysms and infections.
137	8	0.8	309	22 AAG79254	CC	
138	8	0.8	309	22 AAG79255	CC	
139	8	0.8	309	22 AAG79256	CC	
140	8	0.8	309	22 AAG79257	CC	
141	8	0.8	309	22 AAG79258	CC	
142	8	0.8	309	22 AAG79259	CC	
143	8	0.8	309	23 AABP0125	CC	
144	8	0.8	332	21 AAG37725	QY	Sequence 1001 AA;
145	8	0.8	332	21 AAG34478	QY	Query Match 100.0%; Score 1001; DB 22; Length 1001;
146	8	0.8	332	21 AAG34478	QY	Best Local Similarity 100.0%; Pred. No. 0; Matches 1001; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
147	8	0.8	332	21 AAG54121	Db	1 MABPGHSHHLSLARVRGRERLPRWLRLWAGTAQVOTGTGPELHACKSEYHEYTA 60
148	8	0.8	342	10 AAP5203	QY	61 COSTGSRWWRVAVPHPTLCSTSRLDPVKGTEFSFCNAGEFLDMKDOSCKPCEAEGRSLGT 120
149	8	0.8	349	14 AAM32760	Db	61 COSTGSRWWRVAVPHPTLCSTSRLDPVKGTEFSFCNAGEFLDMKDOSCKPCEAEGRSLGT 120
150	8	0.8	353	21 AAG34477	QY	181 VNIKQSTVNPIYYDPSIPEFFVONDQPAQNAQDSRMRKTEKGWEFVSELNRGN 240
					181 VNIKQSTVNPIYYDPSIPEFFVONDQPAQNAQDSRMRKTEKGWEFVSELNRGN 240	
					241 VLYWRTTAFSTWTKVPRVJNIAITGVAYTSECPCPKGTYIAKQGSSFFKLPANSY 300	
					241 VLYWRTTAFSTWTKVPRVJNIAITGVAYTSECPCPKGTYIAKQGSSFFKLPANSY 300	

ALIGNMENTS

RESULT 1

Db	3.01 SNKGGETSCHQCDPDKYSEKGSSCNVRPACTDKDYFTHACDANGETOQMYKWAKPIC	XX	Bruck CEM, Casart J, Coche T, Vinals De Bassols YC;
Qy	3.61 SEDLEGAVKLPASGVKTHCPCPNPGFKTNISTCOPCPGKSYNSGSDCTRCPAGTEPAVG	XX	DR WPI; 2000-664921/64.
Db	3.61 SEDLEGAVKLPASGVKTHCPCPNPGFKTNISTCOPCPGKSYNSGSDCTRCPAGTEPAVG	XX	DR N-PSDB; AAJ95442.
Qy	4.21 FEYKWNNTLPNTMELVLSGINFNEYKGMGTMGEVAGDHITYTAGASDNDMFLITLVPGFR	420	Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases
Db	4.21 FEYKWNNTLPNTMELVLSGINFNEYKGMGTMGEVAGDHITYTAGASDNDMFLITLVPGFR	480	PT PT ovarian and colon carcinoma, and autoimmune diseases
Qy	4.81 PPOSWADTENKEVARITFVETLCSVNCELYFMVGINSRNTPVEIWKGSKGKSYTYI	540	RS Claim 4; Page 54-56; 68pp; English.
Db	4.81 PPOSWADTENKEVARITFVETLCSVNCELYFMVGINSRNTPVEIWKGSKGKSYTYI	540	CC The present sequence comprises the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The invention provides a number of epitopes derived from the protein which can be used as immunogens.
Qy	5.41 IEENTTTSFMAFORTTHEASRKYNDVAKIYSINTVNMGVASYCRCPALEDASVGS	600	CC
Db	5.41 IEENTTTSFMAFORTTHEASRKYNDVAKIYSINTVNMGVASYCRCPALEDASVGS	600	CC
Qy	6.01 SCTSPAGYIIDRSGICHPNNTLKAHQPYGQACVPGCPGTKNNKTHSLCNDCTF	660	CC
Db	6.01 SCTSPAGYIIDRSGICHPNNTLKAHQPYGQACVPGCPGTKNNKTHSLCNDCTF	660	CC
Qy	6.61 SRNTPRTENNFNSALANTVLAGPSFTSKGLKYFHHFTSLCGNQGRKNSVCTNDVTD	720	XX
Db	6.61 SRNTPRTENNFNSALANTVLAGPSFTSKGLKYFHHFTSLCGNQGRKNSVCTNDVTD	720	SQ Sequence 1013 AA;
Qy	7.21 LRIPEBEGSGFSKISTAVCQAVITPPBVGKAGQVSVQPSVSLADRLIGTIDMTLDGITS	780	Query Match 87.6%; Score 877; DB 21; Length 1013;
Db	7.21 LRIPEBEGSGFSKISTAVCQAVITPPBVGKAGQVSVQPSVSLADRLIGTIDMTLDGITS	780	Best Local Similarity 99.9%; Pred. No. 0; Mismatches 0; Indels 0; Gaps 0; Matches 977; Conservative 0;
Qy	7.81 PAELPHIESLGLTPDVFYRSNDVTSOCCSGRSTTRVRCSPQKTVPGSLLPPGTCSDGT	840	Db 1 MAEPGHSHILSARVRGTERTERIPRWRLLWAGTAQVTCGFLHACSESEYVETA 60
Db	7.81 PAELPHIESLGLTPDVFYRSNDVTSOCCSGRSTTRVRCSPQKTVPGSLLPPGTCSDGT	840	Qy 61 CDTGSSWRWRVAVPHPTGLCISLPPVGKTEGCSFCNAGELIDMKDOSCKPCAEGRYSLGT 120
Qy	8.41 CDGCFNFFLWESAAACPLCSVADYHAIVSSCVAGIOTKTYWREPKLCSGSIISLPRQVT	900	Db 61 CDTGSSWRWRVAVPHPTGLCISLPPVGKTEGCSFCNAGELIDMKDOSCKPCAEGRYSLGT 120
Db	8.41 CDGCFNFFLWESAAACPLCSVADYHAIVSSCVAGIOTKTYWREPKLCSGSIISLPRQVT	900	Qy 121 GIRFDWDELPHGEASLSANMELDSAASESTGNTSSKWRGDXIAFNEDECTLIMY 180
Qy	9.01 ICKTIDEWLKGIGSAGCTTAILLTULCYFWKQKQLEYKYSKUWVATLKDCDPAA	960	Db 121 GIRFDWDELPHGEASLSANMELDSAASESTGNTSSKWRGDXIAFNEDECTLIMY 180
Db	9.01 ICKTIDEWLKGIGSAGCTTAILLTULCYFWKQKQLEYKYSKUWVATLKDCDPAA	960	Qy 181 VNLKQSGTVAPEYYPDSITIPEFVQNDQGQNAQDSSRMRKTTKGEWEHTSVELNRGN 240
Qy	9.61 CAIMEGEDVEDDLIFISKHNSIGRSHNLLPQGLMDITQCR 1001		Db 181 VNLKQSGTVAPEYYPDSITIPEFVQNDQGQNAQDSSRMRKTTKGEWEHTSVELNRGN 240
Db	9.61 CAIMEGEDVEDDLIFISKHNSIGRSHNLLPQGLMDITQCR 1001		Qy 241 VLYWRITAFSWTIVKPKVLPVLRNIAITGVATYVSECFPPCKPKGYIADKQGSSFFCKLCPANSY 300
RESULT 2			Db 241 VLYWRITAFSWTIVKPKVLPVLRNIAITGVATYVSECFPPCKPKGYIADKQGSSFFCKLCPANSY 300
AAB26179			Qy 3.01 SNKGGETSCHQCDPDKYSEKGSSCNVRPACTDKDYFTHACDANGETOQMYKWAKPIC 360
ID AAB26179	standard; Protein: 1013 AA.		Db 3.01 SNKGGETSCHQCDPDKYSEKGSSCNVRPACTDKDYFTHACDANGETOQMYKWAKPIC 360
XX			Qy 3.61 SEDLEGAVKLPASGVKTHCPCPNPGFKTNISTCOPCPGKSYNSGSDCTRCPAGTEPAVG
AC			Db 3.61 SEDLEGAVKLPASGVKTHCPCPNPGFKTNISTCOPCPGKSYNSGSDCTRCPAGTEPAVG
XX			Qy 4.21 FEYKWNNTLPNTMELVLSGINFNEYKGMGTMGEVAGDHITYTAGASDNDMFLITLVPGFR
DT	12-FEB-2001 (first entry)		Db 420
XX			Qy 4.21 FEYKWNNTLPNTMELVLSGINFNEYKGMGTMGEVAGDHITYTAGASDNDMFLITLVPGFR
DE	Human CASB619 protein #1.		Db 480
XX			Qy 4.81 PPOSWADTENKEVARITFVETLCSVNCELYFMVGINSRNTPVEIWKGSKGKSYTYI
Human; CASB619; cancer; autoimmune disease; immunogen; vaccine; epitope.			Db 540
XX			Qy 541 IEENTTTSFMAFORTTHEASRKYNDVAKIYSINTVNMGVASYCRCPALEDASVGS
OS	Homo sapiens.		Db 541 IEENTTTSFMAFORTTHEASRKYNDVAKIYSINTVNMGVASYCRCPALEDASVGS
XX			Qy 600
PN	WO200058460-A2.		Db 600
XX			Qy 601 SCTSPAGYIIDRSGICHPNNTLKAHQPYGQACVPGCPGTKNNKTHSLCNDCTF
PD	05-OCT-2000.		Db 660
XX			Qy 661 SRNTPRTENNFNSALANTVLAGPSFTSKGLKYFHHFTSLCGNQGRKNSVCTNDVTD
PF	20-MAR-2000; 2000WO-EP02478.		Db 720
XX			Qy (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
PR	26-MAR-1999; 99GB-0007113.		
XX	25-SEP-1999; 99GB-0022858.		
PA			

Db	661 SRNPPTRTNPFNSALNTVLAGSPSFTSKGLKVFHFTLSLGNQGRKMSVCTDNVTD	XX	PA (GETH) GENENTECH INC.
QY	721 LRPEGESEPSKSTAYVQAVIPPEVYKACVSSQPSVSLADRLLGTTDMFLDGTS	XX	PI Baker KP, Baresini M, Deforge L, Destroyers L, Filvaroff E, Gao W;
Db	721 LRPEGESEPSKSTAYVQAVIPPEVYKACVSSQPSVSLADRLLGTTDMFLDGTS	XX	PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
QY	781 PAELPHLESLGIPVTFYRSNDTQSCSGRSTTIVRCSPOTVPEPSLIPCTCSGDT	XX	PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
Db	781 PAELPHLESLGIPVTFYRSNDTQSCSGRSTTIVRCSPOTVPEPSLIPCTCSGDT	XX	DR WPI; 2001-40281/43.
QY	841 CDGPNFHLMESAAACPIGSVADPHAINSCVAGIQKTYWREPKLCSGGISIPEQRT	XX	DR N-PSDB; AAS21262.
Db	841 CDGPNFHLMESAAACPIGSVADPHAINSCVAGIQKTYWREPKLCSGGISIPEQRT	XX	PT Isolated, secretory and transmembrane PRO polypeptide used to detect other PRO polypeptides, link bioactive molecules to cells expressing PT PRO polypeptides, and detect the presence of mammalian tumours e.g. lung, breast, prostate, cervical
QY	901 ICKTIDFWLKVGISAGTCAILLTVLTCYFWKKNQKLEYKYSKUVMNATLKDCDLPAADS	XX	PT
Db	901 ICKTIDFWLKVGISAGTCAILLTVLTCYFWKKNQKLEYKYSKUVMNATLKDCDLPAADS	XX	PS Claim 12; Fig 38; 813pp; English.
QY	961 CAINEGEDVDDLFITSK 978	XX	CC AAU1217-AAU12446 represent novel human secretory and transmembrane CC PRO polypeptides. The PRO polypeptides are useful to detect other CC PRO polypeptides, to link bioactive molecules to cells expressing CC PRO polypeptides, to modulate biological activities of cells expressing CC PRO polypeptides, and to detect the presence of mammalian lung, colon, CC breast, prostate, rectal, cervical or liver tumours by comparing PRO CC polypeptide expression in a cell sample to that in a control sample. CC Some of the 275 sequences are also useful to stimulate the release of CC tumour necrosis factor-alpha (TNF-alpha) from human blood, the CC proliferation or differentiation of chondrocytes, the proliferation or CC gene expression in pericyte cells, the release of proteoglycans from CC cartilage, the proliferation of inner ear utricular supporting cells or CC of T-lymphocytes, the release of a cytokine from peripheral blood monocytes (PRMCs), or the proliferation of endothelial cells. Some of CC the PRO polypeptides may modulate glucose or free fatty acid uptake by CC skeletal muscle cells or by adipocytes; or inhibit binding of A-peptide CC to factor VITA. The PRO polypeptides can be used in assays to identify CC molecules involved in binding interactions. The polynucleotides encoding CC PRO polypeptides can be used to generate probes, antisense RNA/DNA, CC transgenic or knock out animals and can be used in gene therapy.
OS	XX	XX	Sequence 1013 AA:
XX	WO200140466-A2.	XX	Query Match Best Local Similarity 70.7%; Score 708; DB 22; Length 1013; Matches 708; Similarity 100.0%; Pred. No. 0; Mismatches 0; Indels 0; Gaps 0;
PD	07-JUN-2001.	XX	QY 169 NTBECTATIMAYNLKQSGTVNFEYIPDSSTIFEFVNQNDQCPNADDSRMKTTBKW 228
XX		Db 169 NTBECTATIMAYNLKQSGTVNFEYIPDSSTIFEFVNQNDQCPNADDSRMKTTBKW 228	
PR	01-DEC-1999; 99K0-US28301.	XX	QY 229 EFTISVELNRGGNNLYWRITAFSFWTKTPKPVLAIRNIAITGVAWTSECFPCPKGTYADQG 288
PR	02-DEC-1999; 99K0-US28534.	Db 229 EFTISVELNRGGNNLYWRITAFSFWTKTPKPVLAIRNIAITGVAWTSECFPCPKGTYADQG 288	
PR	02-DEC-1999; 99K0-US28564.	XX	QY 289 SSICKLCPANSYENKGTSCHQCDPDKYSEKSSCNVRPACTDKDXYFTYHACDANGET 348
PR	02-DEC-1999; 99K0-US28565.	Db 289 SSICKLCPANSYENKGTSCHQCDPDKYSEKSSCNVRPACTDKDXYFTYHACDANGET 348	
PR	05-DEC-1999; 99US0-0170262.	XX	QY 349 QLMYKWAQPKICSEDLRGAVKLPASGVKTHCPNCPNPFKINNSTCOPCPGYSNSDC 408
PR	09-DEC-1999; 99US0-030095.	Db 349 QLMYKWAQPKICSEDLRGAVKLPASGVKTHCPNCPNPFKINNSTCOPCPGYSNSDC 408	
PR	16-DEC-1999; 99K0-US30095.	XX	QY 409 TRPAGTEPAGVGEYKWNNTLPINMETVLSGIFTNFEYKGMIGWEVAGDHIVAGASND 468
PR	20-DEC-1999; 99K0-US01911.	Db 409 TRPAGTEPAGVGEYKWNNTLPINMETVLSGIFTNFEYKGMIGWEVAGDHIVAGASND 468	
PR	20-DEC-1999; 99K0-US30999.	XX	Db 469 FMLLTUVPGFRIPOSWMADTENKEVIRTFYFETLCSVNCLYFWMGVNRTNTPEW 528
PR	30-DEC-1999; 99K0-US31243.	XX	QY 529 KGSKGKQSYTTEENTTSFMAQRTTHeASRKYTNDAK1YSINVNTMNGVASYC 588
PR	06-JAN-2000; 2000K0-US00277.	Db 529 KGSKGKQSYTTEENTTSFMAQRTTHeASRKYTNDAK1YSINVNTMNGVASYC 588	
PR	11-FEB-2000; 2000K0-US03565.	XX	PR 02-JUN-2000; 2000K0-US03601.
PR	18-FEB-2000; 2000K0-US04341.	XX	PR 21-MAR-2000; 2000K0-US07377.
PR	18-FEB-2000; 2000K0-US04342.	XX	PR 22-FEB-2000; 2000K0-US04414.
PR	22-FEB-2000; 2000K0-US04414.	XX	PR 24-FEB-2000; 2000K0-US04914.
PR	06-JAN-2000; 2000K0-US00376.	XX	PR 01-MAR-2000; 2000K0-US05004.
PR	05-JAN-2000; 2000K0-US00376.	XX	PR 01-MAR-2000; 2000K0-US05061.
PR	11-FEB-2000; 2000K0-US03565.	XX	PR 18-FEB-2000; 2000K0-US04341.
PR	18-FEB-2000; 2000K0-US04341.	XX	PR 18-FEB-2000; 2000K0-US07377.
PR	18-FEB-2000; 2000K0-US07377.	XX	PR 21-MAR-2000; 2000K0-US07532.
PR	22-FEB-2000; 2000K0-US04414.	XX	PR 30-MAR-2000; 2000K0-US04349.
PR	24-FEB-2000; 2000K0-US04914.	XX	PR 22-MAY-2000; 2000K0-US13705.
PR	01-MAR-2000; 2000K0-US05004.	XX	PR 30-MAY-2000; 2000K0-US14042.
PR	01-MAR-2000; 2000K0-US05061.	XX	PR 02-JUN-2000; 2000K0-US14941.
PR	02-JUN-2000; 2000K0-US15264.	XX	PR 10-NOV-2000; 2000K0-US30873.
PR	10-NOV-2000; 2000K0-US30873.	XX	QY 589 RPCALEASDVGSCTCPAGYYIDRDSGTCCHSCPPNTILKAHQPYGVQACVPCGGPTKNN 648

Db	589	RPCALEASDVSSCTSCPGAYYIDRSGCTCHSCCPNTILKAHQPGVQACPGCPGKTN	648
Qy	649	KIHSCLCYNDCTTSRNTTPTTENVNFSALANTVLAGPSFTSKGLKVFHFTPSLICGNG	708
Db	649	KIHSCLCYNDCTTSRNTTPTTENVNFSALANTVLAGPSFTSKGLKVFHFTPSLICGNG	708
Qy	709	RKNSVCTDNVTLRIPGEGSGSKSTIAYVQAVIIPPEVTGKAGVSSQPSLADRIG	768
Db	709	RKNSVCTDNVTLRIPGEGSGSKSTIAYVQAVIIPPEVTGKAGVSSQPSLADRIG	768
Qy	769	VTIDMTLGDITSPAEFLHESLGPIDVIFYRSNDVTOCSSRSTTIRVRCSPQKTVG	828
Db	769	VTIDMTLGDITSPAEFLHESLGPIDVIFYRSNDVTOCSSRSTTIRVRCSPQKTVG	828
Qy	829	SLLPCTGSDGTCGCFHFLWESAAACPLCSVADYHAIVSSCAGIQ	876
Db	829	SLLPCTGSDGTCGCFHFLWESAAACPLCSVADYHAIVSSCAGIQ	876
RESULT 4			
AAB35328	598	VGSSTSCPAGYIYRDGTHSCCPNTILKAHQPGVQACPGCPGKTNKIHSLCYN	QY
ID	598	VGSSTSCPAGYIYRDGTHSCCPNTILKAHQPGVQACPGCPGKTNKIHSLCYN	QY
AAB35328;	658	CTFSRHTPTPFTNMFNSALANTVLAGPSFTSKGLKVFHFTPSLICGNGQKSVTD	QY
AC	658	CTFSRHTPTPFTNMFNSALANTVLAGPSFTSKGLKVFHFTPSLICGNGQKSVTD	QY
XX	718	VTIDRIPEGSGFSSTIAYVQAVIIPPEVTGKAGVSSQPSLADRIGTTDMID	QY
XX	467	VTIDRIPEGSGFSSTIAYVQAVIIPPEVTGKAGVSSQPSLADRIGTTDMID	QY
DE	407	CTFSRHTPTPFTNMFNSALANTVLAGPSFTSKGLKVFHFTPSLICGNGQKSVTD	QY
DE	778	ITSPAEFLHESLGPIDVIFYRSNDVTOCSSRSTTIRVRCSPQKTVG	QY
XX	527	ITSPAEFLHESLGPIDVIFYRSNDVTOCSSRSTTIRVRCSPQKTVG	QY
OS	838	DGTCIGCNFHFLWESAAACPLCSVADYHAIVSSCAGIQKTYWWRBPKLCSGGGSIPE	QY
XX	587	DGTCIGCNFHFLWESAAACPLCSVADYHAIVSSCAGIQKTYWWRBPKLCSGGGSIPE	QY
XX	898	RVTICKTIDFWLKVGISAGTCAITLTVLTYCFWKQNKQLEYKSYKLVMNATLKDQLP	QY
PR	647	RVTICKTIDFWLKVGISAGTCAITLTVLTYCFWKQNKQLEYKSYKLVMNATLKDQLP	QY
PR	958	ADSCAIMEGVEDDVLIFTSKNSLGRSNHUPRGLIMDUTQCR	1001
PR	707	ADSCAIMEGVEDDVLIFTSKNSLGRSNHUPRGLIMDUTQCR	750
RESULT 5			
DR	AAB83845	AAB83845 standard; Protein; 1013 AA.	
DR	AAB83845	AAB83845 standard; Protein; 1013 AA.	
PT	AC	AAB83845;	
PT	XX	AAB83845;	
PT	DT	23-JUL-2001 (first entry)	
PT	XX	Amino acid sequence of a human protein expressed in tumour cells	
XX	DE	Tumour cell; immunological disease; autoimmune disease; cancer; infection;	
XX	CC	Homo sapiens.	
CC	XX	Key	Location/Qualifiers
CC	XX	Peptide	1..41 /note= "signal peptide"
CC	XX	Domain	42..911
SQ	Sequence	750 AA;	

The present invention provides the protein and coding sequences of the human tumour necrosis factor receptors TR13 and TR14. These sequences are useful in the diagnosis and treatment of many diseases, including cancer, autoimmune diseases, cardiovascular disorders, allergies, neurodegenerative diseases, graft rejection, inflammation, aneurysms and

Query Match: 70.3%; Score: 704; DB: 22; Length: 750;
 Best Local Similarity: 100.0%; Pred. No.: 0;
 Matches: 704; Conservative: 0; Mismatches: 0; Indels: 0; Gaps: 0;

/note= "transmembrane domain"

FT
 XX
 PN WO200131003-A1.
 XX
 PD 03-MAY-2001.
 XX
 PP 30-OCT-2000; 2000WO-FR03032.
 XX
 PR 29-OCT-1999; 99FR-0013629.
 XX
 PA (FABR) FABRE MEDICAMENT SA PIERRE.
 XX
 PT Deineste Y, Magistrelli G, Jeannin P, Bonnefoy J;
 XX
 DR WPI; 2001-328651/34.
 DR N-PSDB; AAF89765.
 XX
 PT New nucleic acid, expressed in tumours and lymphoid tissue is useful for identifying agents for treating tumours and autoimmune disease -
 XX
 PS Claim 9; Page 48-51; 85pp; French.
 XX
 CC The present sequence represents a human protein expressed in tumour cells. The polynucleotide is useful for screening cDNA/genomic DNA banks and for cloning normal DNA, identifying mutant forms of the gene that encodes a human protein, where the mutations are associated with abnormal gene expression, or promoters and regulators of the gene, particularly for diagnosis; for recombinant expression of the derived protein; as probes and primers for detection and amplification; and as antisense therapeutics. The tumour expressed protein is useful for raising specific antibodies to screen agents that modulate its activity, bind to it or interact with it. These agents are potentially useful for treatment or prevention of diseases associated with abnormal expression/activity of the protein, particularly immunological diseases (autoimmune diseases and cancer) or viral, bacterial, fungal or parasitic infections.
 XX
 Sequence 1013 AA:
 SQ

Query Match 67.4%; Score 675; DB 22; Length 1013;
 Best Local Similarity 99.7%; Pred. No. 0; Mismatches 3; Indels 0; Gaps 0;
 Matches 975; Conservative 99.7%; Pred. No. 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 MAEGHSHHLSARYVRGRTERRPRPLRLLWAGTAFQVTOGTGELHACKESYEHYETA 60
 Db 1 MAEGHSHHLSARYVRGRTERRPRPLRLLWAGTAFQVTOGTGELHACKESYEHYETA 60
 QY 61 CDGSTGSRREVAVRPTPGLCTSLPPVKGBCSECNAGERFLMDMDQSCPKPCARGYSIGT 120
 Db 61 CDGSTGSRREVAVRPTPGLCTSLPPVKGBCSECNAGERFLMDMDQSCPKPCARGYSIGT 120
 QY 121 GIRDEWDLPHGASLSANMELDSAESTGNOTSSKAVPRGYIAFTIDECTATMVA 180
 Db 121 GIRDEWDLPHGASLSANMELDSAESTGNOTSSKAVPRGYIAFTIDECTATMVA 180
 QY 181 VNLHQSGTNPFEYYPDSSIFERPVQNDQCPNADDSRMKITEKGWMEFHSLFLRGN 240
 Db 181 VNLHQSGTNPFEYYPDSSIFERPVQNDQCPNADDSRMKITEKGWMEFHSLFLRGN 240
 QY 241 VLYRRTTAFSVWTKVTPKPVLRNAAITGAYTSCFPCPKGTYDQKQSSSFCKLCPANSY 300
 Db 241 VLYRRTTAFSVWTKVTPKPVLRNAAITGAYTSCFPCPKGTYDQKQSSSFCKLCPANSY 300
 QY 301 SNKQRTSCHOCDPKYSKEGSSSNVRPACTDKYFYTACDNGETOLMYKAKPKC 360
 Db 301 SNKQRTSCHOCDPKYSKEGSSSNVRPACTDKYFYTACDNGETOLMYKAKPKC 360
 QY 361 SEDLEGAVKLPASQYKTHCPPCNQFPKINNSTQCPYQPSYSQGSDCTRCPAGEAVG 420
 Db 361 SEDLEGAVKLPASQYKTHCPPCNQFPKINNSTQCPYQPSYSQGSDCTRCPAGEAVG 420
 QY 421 FEYKWWNTLPTNMETVLSGINFEYKGMGTCWAGDHTYTAGASDNDMILTVLPGFR 480
 Db 421 FEYKWWNTLPTNMETVLSGINFEYKGMGTCWAGDHTYTAGASDNDMILTVLPGFR 480
 QY 481 PPGSMADTENKEVARTTFVFFETCSVNCYLYFMVGNSRNTPWTKGSKQSYTI 540
 Db 481 PPQSMADTENKEVARTTFVFFETCSVNCYLYFMVGNSRNTPWTKGSKQSYTI 540
 QY 541 IEENTTSPTWAFTTPEAERKYTDVAKIYSINTVNMGVASTCRPCPCLAEASPGS 600
 Db 541 IEENTTSPTWAFTTPEAERKYTDVAKIYSINTVNMGVASTCRPCPCLAEASPGS 600
 QY 601 SCSCPAGYYIDDSGCHSCPNTIKAHQYQVQCYCPCPGTKONKISLCLYNDCTF 660
 Db 601 SCSCPAGYYIDDSGCHSCPNTIKAHQYQVQCYCPCPGTKONKISLCLYNDCTF 660
 QY 661 SRNTPTRTFNNYNSALANTVLAGGPFTSKGKYFHFTSLCGNQGRKMSVCTDNWD 720
 Db 661 SRNTPTRTFNNYNSALANTVLAGGPFTSKGKYFHFTSLCGNQGRKMSVCTDNWD 720
 QY 721 LRIPEGESGSFSKTSIAYCQAVIPPEVTGKAGVSSQPSVSLADRLLGVTIDWLDGTS 780
 Db 721 LRIPEGESGSFSKTSIAYCQAVIPPEVTGKAGVSSQPSVSLADRLLGVTIDWLDGTS 780
 QY 781 PAELPHLSIGDIDVIFYRNUVQCSGSGTTRVRCSPKTVGSLIUPGTSPT 840
 Db 781 PAELPHLSIGDIDVIFYRNUVQCSGSGTTRVRCSPKTVGSLIUPGTSPT 840
 QY 841 CDGONFHILWESAAACPLCSVADYHATVSSCAGIQKITYWREPKLCSGSLPEORT 900
 Db 841 CDGONFHILWESAAACPLCSVADYHATVSSCAGIQKITYWREPKLCSGSLPEORT 900
 QY 901 ICKTIDFWLKGVSAGTCIAITLTVLTCYFWKKNQKLUBYKXKLMVNTATKQCDLPAADS 960
 Db 901 ICKTIDFWLKGVSAGTCIAITLTVLTCYFWKKNQKLUBYKXKLMVNTATKQCDLPAADS 960
 QY 961 CAAMEGEVEDDILFTSK 978
 Db 961 CAAMEGEVEDDILFTSK 978
 QY 978 CALMEGEDVEDDILFTSK 978

RESULT 6
 AAB83850
 ID AAB83850 standard; Protein; 911 AA.
 XX
 AC AAB83850;
 XX
 DT 23-JUL-2001 (first entry)
 XX
 DE Amino acid sequence of a human protein expressed in tumour cells.
 XX
 KW Tumour cell; immunological disease; autoimmune disease; cancer;
 KW infection.
 XX
 IN Homo sapiens.
 XX
 PN WO200131003-A1.
 XX
 PD 03-MAY-2001.
 XX
 PR 30-OCT-2000; 2000WO-FR03032.
 XX
 PR 29-OCT-1999; 99FR-0013629.
 XX
 PA (FABR) FABRE MEDICAMENT SA PIERRE.
 XX
 PI Deineste Y, Magistrelli G, Jeannin P, Bonnefoy J;
 XX
 DR WPI; 2001-328651/34.
 DR N-PSDB; AAF89774.
 XX
 PT New nucleic acid, expressed in tumours and lymphoid tissue is useful for identifying agents for treating tumours and autoimmune disease -
 XX
 PS Claim 10; Page 60-63; 85pp; French.

XX
 CC The present sequence represents a human protein expressed in tumour
 CC cells. The polynucleotide is useful for screening cDNA/genomic DNA banks
 CC and for cloning isolated DNA; identifying mutant forms of the gene that
 CC encodes a human protein, where the mutations are associated with
 CC abnormal gene expression, or promoters and regulators of the gene,
 CC particularly for diagnosis; for recombinant expression of the derived
 protein; as probes and primers for detection and amplification; and
 CC as antisense therapeutics. The tumour expressed protein is useful for
 CC raising specific antibodies and to screen agents that modulate its
 CC activity, bind to it or interact with it. These agents are potentially
 CC useful for treatment or prevention of diseases associated with abnormal
 CC expression/activity of the protein, particularly immunological diseases
 CC (autoimmune diseases and cancer) or viral, bacterial, fungal or parasitic
 CC infections.
 XX

SQ Sequence 911 AA;

Query Match 66.1%; Score 662; DB 22; Length 911;
 Best Local Similarity 99.8%; Pred. No. 0; Mismatches 2; Indels 0; Gaps 0;

QY

1 MAEPGHSHHLSARVRGTERR1PRLWILLWAGTAQVTOQGPGELACKESBYHYETA 60

Db

61 CDSTGSRMRVAVPHTPGLTSPPVKGTECSFSCNAGEFLDMKQDSCKPCAEGRSILGT 120

Db

61 CDSTGSRMRVAVPHTPGLTSPPVKGTECSFSCNAGEFLDMKQDSCKPCAEGRSILGT 120

QY

121 GIRDPEDELPLPGFFASUSANMELDSAESTGNTCTSSKWVPRGDIYAFNTDECTATMYA 180

Db

121 GIRDPEDELPLPGFFASUSANMELDSAESTGNTCTSSKWVPRGDIYASNTDECTATMYA 180

QY

181 VNLKQSGTVNPFYVYPPSITFEFFVQNDQCPNADDSSRWMKTEKWFHFSVELNRGN 240

Db

181 VNLKQSGTVNPFYVYPPSITFEFFVQNDQCPNADDSSRWMKTEKWFHFSVELNRGN 240

QY

241 VLYWRTTAFASVWIKVPKPVLUVRNIAITGVAYISCFCPCKPGYIADKGSSFFCKLCPANSY 300

Db

241 VLYWRTTAFASVWIKVPKPVLUVRNIAITGVAYISCFCPCKPGYIADKGSSFFCKLCPANSY 300

QY

301 SNGKGETSCHOCPPDKYSEKGSSCNCVERPACTKDYFHTACDANGTQOLMYKWAEPKIC 360

Db

301 SNGKGETSCHOCPPDKYSEKGSSCNCVERPACTKDYFHTACDANGTQOLMYKWAEPKIC 360

QY

361 SEDLEGAVKULPASGVKTHCPCPNPGFFKTNISTCOPPYGYSISNGSDCTRCAGTEAVG 420

Db

361 SEDLEGAVKULPASGVKTHCPCPNPGFFKTNISTCOPPYGYSISNGSDCTRCAGTEAVG 420

QY

421 FEYKWWNLTPTNNTETWLSGNGFENYKGMGTMWEVAGDHIIYTAAGASNDPMLTUVPGFR 480

Db

421 FEYKWWNLTPTNNTETWLSGNGFENYKGMGTMWEVAGDHIIYTAAGASNDPMLTUVPGFR 480

QY

481 PROSMVADTENKEVARITFVFELTSNCYLFMVGNSRNTMPVETWKGSKGKQSYTYI 540

Db

481 PROSMVADTENKEVARITFVFELTSNCYLFMVGNSRNTMPVETWKGSKGKQSYTYI 540

QY

541 IEEETTSPTWAFQRTFHEASRKYNDVAKYISINTNWVAGVASYCRCPALEASDVG 600

Db

541 IEEETTSPTWAFQRTFHEASRKYNDVAKYISINTNWVAGVASYCRCPALEASDVG 600

SQ Sequence 870 AA;

QY

601 SCSGPCAGYIYDRTGSGTCHSCPNPTIKAHQYGVQACVPCPFGTKNNKISLCLYNDCTF 660

Db

601 SCSGPCAGYIYDRTGSGTCHSCPNPTIKAHQYGVQACVPCPFGTKNNKISLCLYNDCTF 660

QY

661 SRSNTPTRFVNFSALANTVTLGGPSFTSKGLYKHFHTLSICGNGRKRMSVCTDNT 720

Db

661 SRSNTPTRFVNFSALANTVTLGGPSFTSKGLYKHFHTLSICGNGRKRMSVCTDNT 720

QY

721 LRPPEGSGSGPSKSTAVCQAVIIPPEFTYKXAGVSSOPVSLADRLIGTTMUDGITS 780

Db

721 LRPPEGSGSGPSKSTAVCQAVIIPPEFTYKXAGVSSOPVSLADRLIGTTMUDGITS 780

QY

841 CCGCNFLWESAAFLCSADY 864

Db

841 CCGCNFLWESAAFLCSADY 864

AC

AAB83051;

XX

DT 23-JUL-2001 (first entry)

XX

DE

Amino acid sequence of a human protein expressed in tumour cells.

XX

Tumour cell; immunological disease; autoimmune disease; cancer; KW infection.

XX

OS

Homo sapiens.

XX

PN WO200131003-A1.

XX

PD 03-MAY-2001.

XX

PF 30-OCT-2000; 2000WO-FR03032.

XX

PR 29-OCT-1999; 99FR-0013629.

XX

PA (FABR) FABRE MEDICAMENT SA PIERRE.

XX

PI Delneste Y, Maistrelli G, Jeannin P, Bonnefoy J;

DR WPI; 2001-328651/34.

XX

N-PSDB; AAF89775.

XX

PT New nucleic acid, expressed in tumours and lymphoid tissue is useful for identifying agents for treating tumours and autoimmune disease -

XX

C10, Page 67-70; 85pp; French.

XX

The present sequence represents a human protein expressed in tumour cells. The polynucleotide is useful for screening cDNA/genomic DNA banks and for cloning isolated DNA; identifying mutant forms of the gene that encodes a human protein, where the mutations are associated with abnormal gene expression, or promoters and regulators of the gene, particularly for diagnosis; for recombinant expression of the derived protein; as probes and primers for detection and amplification; and as antisense therapeutics. The tumour expressed protein is useful for raising specific antibodies and to screen agents that modulate its activity, bind to it or interact with it. These agents are potentially useful for treatment or prevention of diseases associated with abnormal expression/activity of the protein, particularly immunological diseases (autoimmune diseases and cancer) or viral, bacterial, fungal or parasitic infections.

XX

Sequence 870 AA;

QY

62.0%; Score 621; DB 22; Length 870;

Best Local Similarity 99.8%; Pred. No. 0; Mismatches 2; Indels 0; Gaps 0;

Matches 821; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY

42 TGPBLHACKESBYHYETAQVTOQGPGELACKESBYHYETAQVTOQGPGEL 101

Db 1 TGPBLHACKESBYHYETAQVTOQGPGELACKESBYHYETAQVTOQGPGEL 101

QY

102 DMKDQSCPKPCAEGRYSLGTGPFDEDELPLPGFFASUSANMELDSAESTGNTCTSSKWV 161

Db 61 DMKDQSCPKPCAEGRYSLGTGPFDEDELPLPGFFASUSANMELDSAESTGNTCTSSKWV 120

QY	162	RGDYIAFNTECTATIMYAVNLKGSGTVNPEIYDPSS1FEFVQNDQCPNADDSSRM	PA (META-) METAGEN GES GENOMFORSCHUNG MBH.
Db	121	RGDYIAFNTECTATIMYAVNLKGSGTVNPEIYDPSS1FEFVQNDQCPNADDSSRM	XX
QY	222	KTRERKGWEFRSVELNRGNVLYWRITAFSTWTKPCKPKVPLVRNIAITGVAYTSECPCKEG	XX
Db	181	KTRERKGWEFRSVELNRGNVLYWRITAFSTWTKPCKPKVPLVRNIAITGVAYTSECPCKEG	DR
QY	282	TYADKQGSFCKLCPANSYNSKGTSCHQCDPDKYSEKSSCNCVRPACTDKYPTTA	WPI; 1999-591957/51.
Db	241	TYADKQGSFCKLCPANSYNSKGTSCHQCDPDKYSEKSSCNCVRPACTDKYPTTA	N-PSB; AAZ41991.
QY	342	CDANGETOLMYKWKPKKICSEBDLGAVKLPASGKTHCPNPFKPKINSTQCPGCG	281.
Db	301	CDANGETOLMYKWKPKKICSEBDLGAVKLPASGKTHCPNPFKPKINSTQCPGCG	PT
QY	402	YNSQSDCPCPAGEPAPAVFPEYKWNLTPTNEMETVLISINFELKGMTCWEGADHITA	New nucleic acid sequences expressed in uterine cancer tissues, and
Db	361	YNSQSDCPCPAGEPAPAVFPEYKWNLTPTNEMETVLISINFELKGMTCWEGADHITA	derived polypeptides, for treatment of uterine and endometrial cancer
QY	462	AGASDNDFMILTLTUVGFRPQSTPMADTNKEVARITFPEYTCVNCYLFGVGNSRT	and identification of therapeutic agents -
Db	421	AGASDNDFMILTLTUVGFRPQSTPMADTNKEVARITFPEYTCVNCYLFGVGNSRT	XX
QY	522	NTPTEWKSGKGSYTYIIEENTTTSFWAORTFHEASRKYTNDVAKIYNSINTV	Claim 23; Page 287; 444pp; German.
Db	481	NTPTEWKSGKGSYTYIIEENTTTSFWAORTFHEASRKYTNDVAKIYNSINTV	XX
QY	582	NGVASYCRCALESDVGSCSTSPPAGYYIDRSGTCHCPPNITLKAHOPYGVOAC	CC
Db	541	NGVASYCRCALESDVGSCSTSPPAGYYIDRSGTCHCPPNITLKAHOPYGVOAC	This invention describes novel human nucleic acid (cDNA) sequences (A),
QY	642	GPGTKNNKHKSLCNDCTFSRNTPTTFNNFSALANTVTLAGPSFTSKGLKVFHHFTL	CC that are highly expressed in uterine tumour tissue and which have
Db	601	GPGTKNNKHKSLCNDCTFSRNTPTTFNNFSALANTVTLAGPSFTSKGLKVFHHFTL	CC anti-cancer and cytostatic activity. (A) are used (i) to recombinant
QY	702	SLCQNQGRKMSVCTDNVTDLRIPGESEGSKSKTAYCQAVIPPEVYKAGSSQVS	CC expression of polypeptides (B) and (ii) to isolate complete genes. (B)
Db	661	SLCQNQGRKMSVCTDNVTDLRIPGESEGSKSKTAYCQAVIPPEVYKAGSSQVS	CC are used (i) to identify agents suitable for treatment of uterine or
QY	762	LAPRLIGVTTDMTDGITSPAELFLESGLPDIFFYVNSNDTQSCASGRSTTIRVCS	CC endometrial cancer; (ii) directly for treating these forms of cancer
Db	721	LAPRLIGVTTDMTDGITSPAELFLESGLPDIFFYVNSNDTQSCASGRSTTIRVCS	(including expression from gene therapy vectors) and (iii) for
QY	822	POKVPGSLLPGTCSGTDGCFHFLMESAACPLCSVADY	CC generation of specific antibodies. (A) are identified by assembling ESTs
Db	781	POKVPGSLLPGTCSGTDGCFHFLMESAACPLCSVADY	CC (expressed sequence tags) from a particular tissue type before comparison
RESULT 8			CC of expression patterns. This allows a significantly longer fragment of
AY59972			CC the gene to be revealed, so should reduce the number of failures
ID			CC associated with the fact that ESTs from different libraries may represent
XX			CC different parts of the same unknown gene, distorting the estimated
AC			CC frequency of occurrence in a particular tissue. AAI39941-Y60328 represent
AAV59972;			CC protein fragments encoded by the human endometrium tumour cDNA library
DT			CC derived EST fragments represented in AAZ41981-242121.
XX			XX
31-JAN-2000 (first entry)		Sequence 495 AA:	Sequence 495 AA:
XX		Query Match 46.0%; Score 460; DB 20; Length 495;	Query Match 46.0%; Score 460; DB 20; Length 495;
Human endometrium tumour EST encoded protein 32.		Best Local Similarity 100.0%; Pred. No. 0; Mismatches 0; Indels 0; Gaps 0;	Best Local Similarity 100.0%; Pred. No. 0; Mismatches 0; Indels 0; Gaps 0;
KW		Db 519 SRNTPVTWKGSKGSYTYIIEENTTTSFWAORTFHEASRKYTNDVAKIYNSINTV	Db 519 SRNTPVTWKGSKGSYTYIIEENTTTSFWAORTFHEASRKYTNDVAKIYNSINTV
KW		QY 761 1 SRNTPVTWKGSKGSYTYIIEENTTTSFWAORTFHEASRKYTNDVAKIYNSINTV	QY 761 1 SRNTPVTWKGSKGSYTYIIEENTTTSFWAORTFHEASRKYTNDVAKIYNSINTV
DE		Db 661 SLCQNQGRKMSVCTDNVTDLRIPGESEGSKSKTAYCQAVIPPEVYKAGSSQVS	Db 661 SLCQNQGRKMSVCTDNVTDLRIPGESEGSKSKTAYCQAVIPPEVYKAGSSQVS
OS		QY 721 LAPRLIGVTTDMTDGITSPAELFLESGLPDIFFYVNSNDTQSCASGRSTTIRVCS	QY 721 LAPRLIGVTTDMTDGITSPAELFLESGLPDIFFYVNSNDTQSCASGRSTTIRVCS
Homo sapiens.		Db 822 POKVPGSLLPGTCSGTDGCFHFLMESAACPLCSVADY	Db 822 POKVPGSLLPGTCSGTDGCFHFLMESAACPLCSVADY
PN		Db 781 POKVPGSLLPGTCSGTDGCFHFLMESAACPLCSVADY	Db 781 POKVPGSLLPGTCSGTDGCFHFLMESAACPLCSVADY
XX		QY 699 FTSLCQNQGRKMSVCTDNVTDLRIPGESEGSKSKTAYCQAVIPPEVYKAGSSQ	QY 699 FTSLCQNQGRKMSVCTDNVTDLRIPGESEGSKSKTAYCQAVIPPEVYKAGSSQ
DE1981798-A1.		Db 181 FTSLCQNQGRKMSVCTDNVTDLRIPGESEGSKSKTAYCQAVIPPEVYKAGSSQ	Db 181 FTSLCQNQGRKMSVCTDNVTDLRIPGESEGSKSKTAYCQAVIPPEVYKAGSSQ
21-OCT-1999.		QY 759 PVS1ADR1GVTDMTDGITSPAELFLESGLPDIFFYVNSNDTQSCASGRSTTIRV	QY 759 PVS1ADR1GVTDMTDGITSPAELFLESGLPDIFFYVNSNDTQSCASGRSTTIRV
XX		Db 241 PVS1ADR1GVTDMTDGITSPAELFLESGLPDIFFYVNSNDTQSCASGRSTTIRV	Db 241 PVS1ADR1GVTDMTDGITSPAELFLESGLPDIFFYVNSNDTQSCASGRSTTIRV
17-APR-1998;		QY 819 RCSQPKTVPGSLLPGTCSGTDGCFHFLMESAACPLCSVADYHAIVSSCVQIQT	QY 819 RCSQPKTVPGSLLPGTCSGTDGCFHFLMESAACPLCSVADYHAIVSSCVQIQT
XX		Db 301 RCSQPKTVPGSLLPGTCSGTDGCFHFLMESAACPLCSVADYHAIVSSCVQIQT	Db 301 RCSQPKTVPGSLLPGTCSGTDGCFHFLMESAACPLCSVADYHAIVSSCVQIQT
Endometrium; human; tumour; cancer; anticancer; cytostatic; EST:		QY 879 TYWREPKLGGGSLPQRQVICKIDFWLKIGISAGTCAILLTWTCFWKKNQLE	QY 879 TYWREPKLGGGSLPQRQVICKIDFWLKIGISAGTCAILLTWTCFWKKNQLE
XX		Db 361 TYWREPKLGGGSLPQRVICKIDFWLKIGISAGTCAILLTWTCFWKKNQLE	Db 361 TYWREPKLGGGSLPQRVICKIDFWLKIGISAGTCAILLTWTCFWKKNQLE
OS		QY 939 YKTSKLVMNATKDCDLPADSCAIMEGEDVDDLIFSK	QY 939 YKTSKLVMNATKDCDLPADSCAIMEGEDVDDLIFSK
Hom sapiens.		Db 421 YKTSKLVMNATKDCDLPADSCAIMEGEDVDDLIFSK	Db 421 YKTSKLVMNATKDCDLPADSCAIMEGEDVDDLIFSK
RESULT 9			
XX		AAB83853	
XX		ID AAB83853	
XX		standard; Protein; 383 AA.	

AC	AAB83853;	XX	XX
DT	23-JUL-2001 (first entry)	Db	334 FMLILTVPGFRPPQSWADTENKEVARITVFVETLCSVNCELYFMMGVN 383
XX		RESULT 10	
DE	Amino acid sequence of a human protein expressed in tumour cells.	AAB85768	
KW	Tumour cell; immunological disease; autoimmune disease; cancer; infection.	ID AAB85768 standard; Protein; 372 AA.	
KW		XX	
OS	Homo sapiens.	AC AAB85768;	
XX		XX	
PN	WO200131003-A1..	DT 29-OCT-2001 (first entry)	
XX		XX	
PD	03-MAY-2001.	DE Human seven-transmembrane protein 50288 sequence.	
XX		XX	
PF	30-OCT-2000; 2000WO-FR03032.	KW Seven-transmembrane protein; G-protein coupled receptor; GPCR; human; 17724; 50288; 31945; antiinflammatory; antiulcer; cytotactic; virucide; hepatotropic; immunosuppressive; gynecological; neuroprotective; anti-HIV; immunostimulant; dermatological; antiatherosclerotic; cardiant; antiangiadic; antiParkinsonian; nephrotopic; antiHIV; hemotactic; cerebroprotective; osteopathic; analgesic; gene therapy; nootropic.	
XX		KW anti-HIV; immunostimulant; dermatological; antiatherosclerotic; cardiant; antiangiadic; antiParkinsonian; nephrotopic; antiHIV; hemotactic; cerebroprotective; osteopathic; analgesic; gene therapy; nootropic.	
PR	29-OCT-1999; 99FR-0013629.	XX	
XX		XX	
PA	(FABR) FABRE MEDICAMENT SA PIERRE.	DR DR; WPI; 2001-328651/34.	
XX		XX	
PR	Delnestre Y, Magistrelli G, Jeannin P, Bonnefoy J;	PS N-PSDB; AAF9777.	
XX		XX	
DR	WPI; 2001-328651/34.	PN PN	
XX		XX	
PT	New nucleic acid, expressed in tumours and lymphoid tissue is useful for identifying agents for treating tumours and autoimmune disease	PD 16-AUG-2001.	
XX		XX	
PT	Claim 10; Page 74-75; 85pp; French.	PI 12-FEB-2001; 2001WO-US04536.	
PS		PR 11-FEB-2000; 2000US-0182061.	
XX		XX	
CC	The present sequence represents a human protein expressed in tumour cells. The polynucleotide is useful for screening cDNA/genomic DNA banks and for cloning isolated DNA; identifying mutant forms of the gene that encodes a human protein, where the mutations are associated with abnormal gene expression, or promoters and regulators of the gene, particularly for diagnosis; for recombinant expression of the derived protein; as probes and primers for detection and amplification, and as antisense therapeutics. The tumour expressed protein is useful for raising specific antibodies and to screen agents that modulate its activity, bind to it or interact with it. These agents are potentially useful for treatment or prevention of diseases associated with abnormal expression/activity of the protein, particularly immunological diseases (autoimmune diseases and cancer) or viral, bacterial, fungal or parasitic infections.	PA (MILL-) MILLENNIUM PHARM INC.	
CC		XX	
CC		PI Glucksmann MA, Silos-Santiago I;	
CC		DR WPI; 2001-514670/56.	
CC		DR N-PSDB; AAH6195, AAH76196.	
CC		XX	
CC	New seven-transmembrane protein/G-protein coupled receptor polypeptides and polynucleotides for diagnosing, treating seven-transmembrane protein/receptor-related disorders and to identify modulators of therapeutic use	PT New seven-transmembrane protein/G-protein coupled receptor polypeptides and polynucleotides for diagnosing, treating seven-transmembrane protein/receptor-related disorders and to identify modulators of therapeutic use	
CC		XX	
CC	PT therapeutic use	PS Claim 8; Page 139-141; 144pp; English.	
CC		XX	
CC	The invention provides isolated seven-transmembrane protein/G-protein coupled receptor polypeptides selected from 17724, 50288, 31945 proteins. The polypeptides can be expressed by standard recombinant methodology. Modulators of the polypeptides can be identified using a competition binding assay or an assay for receptor-mediated signal transduction. The polypeptides and polynucleotides are useful as reagents or targets in seven-transmembrane protein/receptor assays applicable to treatment and diagnosis of seven-transmembrane protein/receptor-mediated disorders (see AAB76191 for a detailed description of the various disorders that can be treated or diagnosed using the polypeptides). The polynucleotides that are useful to detect mutations in genes and gene expression products such as mRNA, as antisense constructs to control gene expression and for chromosome identification. The present sequence represents the human seven transmembrane protein 50288 sequence.	CC	
CC		XX	
SQ	Sequence 383 AA:	SO Sequence 372 AA:	
Query Match	24.9%; Score 249; DB 22; Length 383;	Query Match	21.6%; Score 216; DB 22; Length 372;
Best Local Similarity	99.7%; Pred. No. 1 5e-253;	Best Local Similarity	99.6%; Pred. No. 9 8e-219;
Matches	0; Mismatches 1; Indels 0; Gaps 0;	Matches	0; Mismatches 1; Indels 0; Gaps 0;
QY	169 NTDECTATLMYAVNLKGSTQTFYYFDDSSITFEFVQNDQCPQRNADDSRMKTEKGW 228	QY 229 EPHSVELNRLGNNVLYMRTTAFSWMVTKVPKVLPVNLTAITGVAYTSICFPCKPGTYADKG 288	QY 409 TRCPAGTEPAGFPEYKWNNTLPNMEITVLSGINPEYKGMCWEGADHITAGASND 468
Db	34 NIDECTATLMTAVNLKGSTQTFYYFDDSSITFEFVQNDQCPQRNADDSRMKTEKGW 93	Db 94 EPHSVELNRLGNNVLYMRTTAFSWMVTKVPKVLPVNLTAITGVAYTSICFPCKPGTYADKG 153	Db 274 TRCPAGTEPAGFPEYKWNNTLPNMEITVLSGINPEYKGMCWEGADHITAGASND 333
QY	289 SSFKLCPANSISNKGETSCHOCDPKYSEKQSSSCNVRPACTKDQFYTHACDANGER 348	QY 289 SSFKLCPANSISNKGETSCHOCDPKYSEKQSSSCNVRPACTKDQFYTHACDANGER 348	QY 469 FMLILTVPGFRPPQSWADTENKEVARITVFVETLCSVNCELYFMMGVN 518
Db	154 SSFKLCPANSISNKGETSCHOCDPKYSEKQSSSCNVRPACTKDQFYTHACDANGER 213	Db 154 SSFKLCPANSISNKGETSCHOCDPKYSEKQSSSCNVRPACTKDQFYTHACDANGER 213	Db 61 CDSTGSRWRVAPHTGPGTSLDPKGTEFSFCAGFEDMKDOSCKPAGBRGLSGT 120
QY	349 QLMYKWKPKCISSELDLGAVALKUPASGVYKTHCPCPNCPRFKTINSTCOPCPGYSNSDC 408	QY 349 QLMYKWKPKCISSELDLGAVALKUPASGVYKTHCPCPNCPRFKTINSTCOPCPGYSNSDC 408	QY 61 MAEPGSHHHLARVRRTERRIPRRLILLWAGTAQVTOGNGPELHACKSEYHEYTA 60
Db	214 QLMYKWKPKCISSELDLGAVALKUPASGVYKTHCPCPNCPRFKTINSTCOPCPGYSNSDC 273	Db 214 QLMYKWKPKCISSELDLGAVALKUPASGVYKTHCPCPNCPRFKTINSTCOPCPGYSNSDC 273	Db 1 MAEPGSHHHLARVRRTERRIPRRLILLWAGTAQVTOGNGPELHACKSEYHEYTA 60
QY	409 TRCPAGTEPAGFPEYKWNNTLPNMEITVLSGINPEYKGMCWEGADHITAGASND 468	QY 409 TRCPAGTEPAGFPEYKWNNTLPNMEITVLSGINPEYKGMCWEGADHITAGASND 468	QY 61 MAEPGSHHHLARVRRTERRIPRRLILLWAGTAQVTOGNGPELHACKSEYHEYTA 60
Db	274 TRCPAGTEPAGFPEYKWNNTLPNMEITVLSGINPEYKGMCWEGADHITAGASND 333	Db 274 TRCPAGTEPAGFPEYKWNNTLPNMEITVLSGINPEYKGMCWEGADHITAGASND 333	Db 61 CDSTGSRWRVAPHTGPGTSLDPKGTEFSFCAGFEDMKDOSCKPAGBRGLSGT 120

CC AAB54007 represent sequences used in the exemplification of the present invention.

CC XX

SQ Sequence 208 AA;

Query Match 15.0%; Score 150; DB 21; length 208;

Best Local Similarity 100.0%; Pred. No. 2; e-149;

Matches 150; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Db 181 VNLKQSGTNFEEYYPDSIIFEFVQNDQCPNADDSSRWMKTEKGWNEFHSELNRNN 240

QY 241 VLYWRITTAESWVWVPUKPUVLYRVAITGAYTBCFPCKPGTYADQGSSFECKLCPANSY 300

Db 241 VLYWRITTAESWVWVPUKPUVLYRVAITGAYTBCFPCKPGTYADQGSSFECKLCPANSY 300

QY 301 SNKGKETSCHQCDPKYS 317

Db 301 SNKGKETSCHQCDPKYS 317

RESULT 11

AAB5342

ID AAB5342 standard; Protein; 208 AA.

XX

AC AAB5342;

XX DT 09-MAR-2001 (first entry)

XX DE Human colon cancer antigen protein sequence SEQ ID NO:982.

XX KW Human; colon cancer; colon cancer antigen; diagnosis; detection; identification; cytosolic; cardioactive; neuroprotective; vulnerary; immunomodulatory; muscular; gynaecological; gastrointestinal; nephrotoxic; anti-infective; antibacterial; gene therapy; wound; neural disorder; immune system disorder; muscular disorder; reproductive disorder; gastrointestinal disorder; renal disorder; infectious disease; cardiovascular disorder.

XX OS Homo sapiens.

XX PN WO200053351-A1.

XX PD 21-SEP-2000.

XX PF 08-MAR-2000; 2000WO-US05883.

XX PR 12-MAR-1999; 99US-0124270.

XX PA (HUMA-) HUMAN GENOME SCI INC.

XX PI Rosen CA, Ruben SM;

XX DR WPI; 2000-587534/55.

XX DR N-FSDB; AAC98199.

XX PT Colon cancer associated gene sequences, referred to as colon cancer antigens, useful for the treatment, prevention, and diagnosis of colon disorders, such as colon cancer -

XX PS Claim 11; Page 155; 2104pp; English.

XX AAC7991 to AAC98753 encode the human colon cancer associated proteins, called human colon cancer antigens, given in AAB5334 to AAB54006. The human colon cancer antigens can have cytosolic, cardioactive, muscular, neuroprotective, immunomodulatory, gynaecological, gastrointestinal, vulnerary, nephrotoxic, anti-infective and antibacterial activities, and can be used in gene therapy. The colon cancer antigen polynucleotides, proteins and antibodies to the proteins are useful for the prevention, treatment and diagnosis of colon disorders, such as colon cancer. The polynucleotides may be used in diagnostics and research, such as for chromosome identification, and as hybridisation probes. The proteins may also be used to prevent diseases such as neural disorders, immune system disorders, muscular disorders, reproductive disorders, gastrointestinal disorders, wounds, renal disorders, infectious diseases, and cardiovascular disorders. AAC98764 to AAC98772 and

CC CC

CC AAB54007 represent sequences used in the exemplification of the present invention.

CC XX

SQ Sequence 208 AA;

Query Match 15.0%; Score 150; DB 21; length 208;

Best Local Similarity 100.0%; Pred. No. 2; e-149;

Matches 150; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Db 181 VNLKQSGTNFEEYYPDSIIFEFVQNDQCPNADDSSRWMKTEKGWNEFHSELNRNN 240

QY 241 VLYWRITTAESWVWVPUKPUVLYRVAITGAYTBCFPCKPGTYADQGSSFECKLCPANSY 300

Db 241 VLYWRITTAESWVWVPUKPUVLYRVAITGAYTBCFPCKPGTYADQGSSFECKLCPANSY 300

QY 301 SNKGKETSCHQCDPKYS 317

Db 301 SNKGKETSCHQCDPKYS 317

RESULT 12

AAB83849

ID AAB83849 standard; peptide; 147 AA.

XX AC AAB83849;

XX DT 23-JUL-2001 (first entry)

XX DE Peptide fragment of a human protein expressed in tumour cells.

XX KW Tumour cell; immunological disease; autoimmune disease; cancer; infection.

XX OS Homo sapiens.

XX PN WO200131003-A1.

XX PD 03-MAY-2001.

XX PR 30-OCT-2000; 2000WO-FR03032.

XX PR 29-OCT-1999; 99FR-0013629.

XX PA (FABR) FABRE MEDICAMENT SA PIERRE.

XX PI Delneste Y, Magistrelli G, Jeannin P, Bonnefoy J;

XX DR WPI; 2001-328651/34.

XX DR N-FSDB; AAF89769.

XX PT New nucleic acid, expressed in tumours and lymphoid tissue is useful for identifying agents for treating tumours and autoimmune disease -

XX PT

XX PS Claim 10; Page 54-55; 85pp; French.

XX CC AAB83846-49 represent fragments of a human protein expressed in tumour cells. The polynucleotide is useful for screening cDNA/genomic DNA banks and for cloning isolated DNA; identifying mutant forms of the gene that encodes a human protein, where the mutations are associated with abnormal gene expression, or promoters and regulators of the gene, particularly for diagnosis; for recombinant expression of the derived protein; as probes and primers for detection and amplification; and as antisense therapeutics. The tumour expressed protein is useful for raising specific antibodies and to screen agents that modulate its activity, bind to it or interact with it. These agents are potentially useful for treatment or prevention of diseases associated with abnormal expression/activity of the protein, particularly immunological diseases (autoimmune diseases and cancer) or viral, bacterial, fungal or parasitic infections.

XX SQ Sequence 147 AA;

Query Match	14.7%	Score 147;	DB 22;	Length 147;	YSGNSDCTRCPAGTEPAGVGEYKWNNTIPTNMENTVLSQINFEYKGMGWEVAGDHITA	461	
Best Local Similarity	100.0%	Pred. No. 2.9e-146;					
Mismatches	147;	Conservative	0;	Mismatches	0;		
Indels	0;			Gaps	0;		
o							
654 CYNDCTFSRNTPIRTFNMFNSALANTVLAGSISGFTSKQLKLYFFHFTSLCGNGRKNV	713	QY	462 AGASDNDFMLTUVPGPRPPOSMDAENKEYARITYVFEETICSVNGEFLYFMGN	518			
61 CTDNVTDLRIPEGSGFSKSTIAYVCQAVIIPPEVTGKAGVASSQPVSLADRLIGVTD	120	Db	153 AGASDNDMILTLVPGPRPPOSMDAENKEYARITYVFEETICSVNGEFLYFMGN	209			
o							
1 CYNDCTFSRNTPIRTFNMFNSALANTVLAGSISGFTSKQLKLYFFHFTSLCGNGRKNV	60	QY	93 YSGNSDCTRCPAGTEPAGVGEYKWNNTIPTNMENTVLSQINFEYKGMGWEVAGDHITA	152			
714 CTDNVTDLRIPEGSGFSKSTIAYVCQAVIIPPEVTGKAGVASSQPVSLADRLIGVTD	773	Db	121 TLDGITSPAELFEHLESLGIPDVIFFYR	147			
o							
774 TLDGITSPAELFEHLESLGIPDVIFFYR	800	QY	462 AGASDNDFMLTUVPGPRPPOSMDAENKEYARITYVFEETICSVNGEFLYFMGN	518			
o							
121 TLDGITSPAELFEHLESLGIPDVIFFYR	147	Db	153 AGASDNDMILTLVPGPRPPOSMDAENKEYARITYVFEETICSVNGEFLYFMGN	209			
o							
RESULT 13							
AB83852							
AB83852 standard; Protein; 209 AA.							
o							
AB83852;							
AB83852;							
23-JUL-2001 (first entry)							
EE							
Amino acid sequence of a human protein expressed in tumour cells.							
Tumour cell; immunological disease; autoimmune disease; cancer; infection.							
EE							
Homo sapiens.							
W0200131003-A1.							
EE							
03-MAY-2001.							
EE							
30-OCT-2000; 2000WO-FR03032.							
EE							
29-OCT-1999; 99FR-0013629.							
EE							
(FABR) FABRE MEDICAMENT SA PIERRE.							
EE							
Delneste Y, Magistrelli G, Jeannin P, Bonnefoy J;							
EE							
WPI; 2001-328651/34.							
EE							
N-PSDB; AAF89776.							
EE							
The present sequence represents a human protein expressed in tumour cells. The polynucleotide is useful for screening cDNA/genomic DNA banks and for cloning isolated DNA; identifying mutant forms of the gene that encodes a human protein, where the mutations are associated with abnormal gene expression, or promoters and regulators of the gene, particularly for diagnosis; for recombinant expression of the derived protein, as probes and primers for detection and amplification; and as antisense therapeutics. The tumour expressed protein is useful for raising specific antibodies and to screen agents that modulate its activity, bind to it or interact with it. These agents are potentially useful for treatment or prevention of diseases associated with abnormal expression/activity of the protein, particularly immunological diseases (autoimmune diseases and cancer) or viral, bacterial, fungal or parasitic infections.							
EE							
Sequence 209 AA;							
EE							
Query Match	9.6%	Score 96;	DB 21;	Length 105;			
Best Local Similarity	100.0%	Pred. No. 1.4e-92;					
Mismatches	96;	Conservative	0;	Mismatches	0;		
Indels	0;			Gaps	0;		
o							
QY	864 YHAIIVSSCAGQKTTWREPKLCSGGISLPEQRVICKTIDFWLKGISAGTCAILL	923	QY	924 TWTCTYFWKKNQKLEYKSKLUNATKCOLPAAD	959	QY	
o							
1 YHAIIVSSCAGQKTTWREPKLCSGGISLPEQRVICKTIDFWLKGISAGTCAILL	60	Db	61 TWTCTYFWKKNQKLEYKSKLUNATKCOLPAAD	96	Db	61 TWTCTYFWKKNQKLEYKSKLUNATKCOLPAAD	96

AC AAB83847;
 XX DT 23-JUL-2001 (first entry)
 XX DE Peptide fragment of a human protein expressed in tumour cells.
 XX KW Tumour cell; immunological disease; autoimmune disease; cancer;
 XX infection.
 OS Homo sapiens.
 XX PN WO200131003-A1.
 XX PR 03-MAY-2001.
 XX PT 30-OCT-2000; 2000WO-FR03032.
 XX PA 29-OCT-1999; 99FR-0013629.
 XX PA (FABR) FABRE MEDICAMENT SA PIERRE.
 XX PT Delneste Y, Magistrelli G, Jeannin P, Bonnefoy J;
 XX PR WPI; 2001-328651/34.
 XX DR N-PSDB; AAF89768.
 XX PR 03-MAY-2001.
 XX PT 30-OCT-2000; 2000WO-FR03032.
 XX PS 29-OCT-1999; 99FR-0013629.
 XX PA (FABR) FABRE MEDICAMENT SA PIERRE.
 XX PI Delneste Y, Magistrelli G, Jeannin P, Bonnefoy J;
 XX PR WPI; 2001-328651/34.
 XX DR N-PSDB; AAF89767.
 XX PR 03-MAY-2001.
 XX PT 30-OCT-2000; 2000WO-FR03032.
 XX PS 29-OCT-1999; 99FR-0013629.
 XX CC AAB83846-49 represent fragments of a human protein expressed in tumour cells. The polynucleotide is useful for screening cDNA/genomic DNA banks and for cloning isolated DNA; identifying mutant forms of the gene that encodes a human protein, where the mutations are associated with abnormal gene expression or promoters and regulators of the gene, particularly for diagnosis; for recombinant expression of the derived protein, as probes and primers for detection and amplification; and as antisense therapeutics. The tumour expressed protein is useful for raising specific antibodies and to screen agents that modulate its activity, bind to it or interact with it. These agents are potentially useful for treatment or prevention of diseases associated with abnormal expression/activity of the protein, particularly immunological diseases (autoimmune diseases and cancer) or viral, bacterial, fungal or parasitic infections.
 XX CC AAB83846-49 represent fragments of a human protein expressed in tumour cells. The polynucleotide is useful for screening cDNA/genomic DNA banks and for cloning isolated DNA; identifying mutant forms of the gene that encodes a human protein, where the mutations are associated with abnormal gene expression, or promoters and regulators of the gene, particularly for diagnosis; for recombinant expression of the derived protein, as probes and primers for detection and amplification; and as antisense therapeutics. The tumour expressed protein is useful for raising specific antibodies and to screen agents that modulate its activity, bind to it or interact with it. These agents are potentially useful for treatment or prevention of diseases associated with abnormal expression/activity of the protein, particularly immunological diseases (autoimmune diseases and cancer) or viral, bacterial, fungal or parasitic infections.
 XX SQ Sequence 52 AA;
 XX ID Query Match 5.6%; Score 56; DB 22; Length 56;
 XX Best Local Similarity 100.0%; Pred. No. 1.4e-50; Mismatches 0; Indels 0; Gaps 0;
 XX Matches 56; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 XX QY 304 GFTSSQCDPKYSKKGSSCNTRPACTDKYFTHTACDANGTQIIMWKAKEPKI 359
 XX DB 1 LTIVWPGRRPQSMWATENKEVARITFVFTLCSNCYELFMVGNSRTNT 523
 XX ID 1 LTIVWPGRRPQSMWATENKEVARITFVFTLCSNCYELFMVGNSRTNT 52
 XX SQ Sequence 52 AA;
 XX ID Query Match 5.2%; Score 52; DB 22; Length 52;
 XX AC Best Local Similarity 100.0%; Pred. No. 2.1e-46; Mismatches 0; Indels 0; Gaps 0;
 XX AC Matches 52; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 XX DT 23-JUL-2001 (first entry)
 XX PR Peptide fragment of a human protein expressed in tumour cells.
 XX KW Tumour cell; immunological disease; autoimmune disease; cancer;
 XX infection.
 OS Homo sapiens.
 XX PN WO200131003-A1.
 XX PR 03-MAY-2001.
 XX PT 30-OCT-2000; 2000WO-FR03032.
 XX PR 29-OCT-1999; 99FR-0013629.
 XX PA (FABR) FABRE MEDICAMENT SA PIERRE.
 XX PI Delneste Y, Magistrelli G, Jeannin P, Bonnefoy J;
 XX PR WPI; 2001-328651/34.
 XX DR N-PSDB; AAF89766.

RESULT 16
 AAB83848
 ID AAB83848 standard; peptide; 52 AA.
 XX AC AAB83848;
 XX DT 23-JUL-2001 (first entry)
 XX DE Peptide fragment of a human protein expressed in tumour cells.
 XX KW Tumour cell; immunological disease; autoimmune disease; cancer;
 XX infection.
 OS Homo sapiens.
 XX PN WO200131003-A1.

PT New nucleic acid, expressed in tumours and lymphoid tissue is useful for identifying agents for treating tumours and autoimmune disease -
 XX
 PS
 CC AAB83846-49 represent fragments of a human protein expressed in tumour cells. The polynucleotide is useful for screening cDNA/genomic DNA bank and for cloning isolated DNA; identifying mutant forms of the gene that encodes a human protein, where the mutations are associated with abnormal gene expression, or promoters and regulators of the gene, particularly for diagnosis; for recombinant expression of the derived protein; as probes and primers for detection and amplification; and as antisense therapeutics. The tumour expressed protein is useful for raising specific antibodies and to screen agents that modulate its activity, bind to it or interact with it. These agents are potentially useful for treatment or prevention of diseases associated with abnormal expression/activity of the protein, particularly immunological diseases (autoimmune disease and cancer) or viral, bacterial, fungal or parasitic infections.
 XX
 SQ Sequence 45 AA:
 Query Match 4.5%; Score 45; DB 22; Length 45;
 Best Local Similarity 100.0%; Pred. No. 4.5e-39;
 Matches 45; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 Qy 1 MAEPGSHSHLSARVGRGERRPRWLWLLWAGTAFOVQGTRGP 45
 Db 1 MAEPGSHSHLSARVGRGERRPRWLWLLWAGTAFOVQGTRGP 45
 RESULT 18
 AAY12274
 ID AAY12274 standard; Protein; 150 AA.
 XX
 AC AAY12274;
 XX
 DT 17-JUN-1999 (first entry)
 XX
 DE Human 5' EST secreted protein SBQ ID NO:305.
 KW Human; secreted protein; EST; expressed sequence tag; diagnosis;
 KW forensic; gene therapy; chromosome mapping; signal peptide;
 upstream regulatory sequence; cytokine activity; cell proliferation;
 differentiation; haemopoiesis regulation; tissue growth regulation;
 reproductive hormone regulation; chemotactic; chemokinetic; haemostatic;
 thrombolytic; anti-inflammatory; tumour inhibition.
 KW Homo sapiens.
 OS Homo sapiens.
 XX
 PN WO9906548-A2.
 XX
 PD 11-FEB-1999.
 XX
 PP 31-JUL-1998; 98WO-1B01222.
 XX
 PR 01-AUG-1997; 97US-0905135.
 XX
 PA (GEST) GENSET.
 XX
 PI Ducleir A, Dumas Milne Edwards J, Lacroix B;
 XX
 DR WPI; 1999-153778/13.
 XX
 N-PSDB; AX41107.
 XX
 PT New nucleic acids encoding human secreted proteins - obtained from
 PT DNA libraries prepared from e.g. liver, ovary, brain, prostate,
 PT kidney, lung, umbilical cord, placenta and colon tissue
 XX
 Claim 27; Page 655-656; 824pp; English.
 XX
 PS AX41094 to AX41347 represent 5' expressed sequence tags (ESTs) for
 PT human secreted proteins, and encode the proteins given in AAY12261 to
 PT AAY12514, respectively. The proteins given represent the signal peptide
 XX
 N-PSDB; AX41107.
 XX

XX
 DE TR16-short "receptor protein.
 XX PR 30-JUN-2000; 2000US-0608408.
 KW PR 03-AUG-2000; 2000US-0632366.
 KW PR 21-SEP-2000; 2000US-0234387.
 KW PR 27-SEP-2000; 2000US-0236359.
 OS PR 04-OCT-2000; 2000GB-0024263.
 Unidentified.
 XX PA (MOLE-) MOLECULAR DYNAMICS INC.
 XX PI Penn SG, Hanzel DK, Chen W, Rank DR;
 XX DR WPI; 2001-483447/52.
 XX Human genome-derived single exon nucleic acid probes useful for
 PT analyzing gene expression in human fetal liver.
 XX PS Claim 27; SEQ ID NO 32316; 639pp + sequence listing; English.
 XX The invention relates to a single exon nucleic acid probe for
 CC measuring human gene expression in a sample derived from human foetal
 CC liver. The single exon nucleic acid probes may be used for predicting,
 CC measuring and displaying gene expression in samples derived from human
 CC fetal liver. The present sequence is a peptide encoded by a single exon
 CC nucleic acid probe of the invention.
 CC Note: The sequence data for this patent did not form part of the
 CC printed specification, but was obtained in electronic format directly
 CC from WIPO at frp.wipo.int/pub/published_pct_sequences.
 XX Sequence 50 AA;
 PT New nucleic acid molecule encoding a TR16 tumor necrosis factor
 PT receptor polypeptide, useful for the diagnosis and treatment of cancer,
 PT autoimmune disorders and cardiovascular disease.
 XX
 PS Claim 1; Fig 1; 286pp; English.
 XX The present invention relates to a TR16 receptor (tumour necrosis
 CC factor receptor superfamily). The invention is useful treating
 CC diseases and disorders associated with the inhibited or increased
 CC apoptosis. In particular inflammatory diseases, cancers, immune and
 CC neurodegenerative disorders may be treated.
 XX Sequence 963 AA;
 SQ
 Query Match 1.5%; Score 15; DB 22; Length 963;
 Best Local Similarity 100.0%; Pred. No. 3.3e-06;
 Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 AC ABB24346 standard; Protein; 50 AA.
 XX
 AC ABB24346;
 XX DT 23-JAN-2002 (first entry)
 DB Protein #6345 encoded by probe for measuring heart cell gene expression.
 XX Human; gene expression; heart; microarray; vascular system;
 KW cardiovascular disease; hypertension; cardiac arrhythmia;
 KW congenital heart disease.
 XX OS Homo sapiens.
 XX PN WO200157274-A2.
 XX PD 09-AUG-2001.
 XX PF 30-JAN-2001; 2001WO-US00666.
 XX PR 04-FEB-2000; 2000US-0180312.
 PR 26-MAY-2000; 2000US-0207456.
 PR 30-JUN-2000; 2000US-0608408.
 PR 03-AUG-2000; 2000US-0632366.
 PR 21-SEP-2000; 2000US-0234387.
 PR 27-SEP-2000; 2000US-0236359.
 PR 04-OCT-2000; 2000GB-0024263.
 XX PA (MOLE-) MOLECULAR DYNAMICS INC.
 XX PI Penn SG, Hanzel DK, Chen W, Rank DR;
 XX DR WPI; 2001-488899/53.

XX
XX Single exon nucleic acid probes for analyzing gene expression in human
PT
PT hearts -
XX
XX
PS
PS Claim 15; SEQ ID NO 26116; 530pp; English.

CC
CC The present invention relates to single exon nucleic acid probes for
CC measuring human gene expression in a sample derived from human heart (see
CC ABA21535; ABA41305). The present sequence is a protein encoded by one such
CC probe. The probes may be used for predicting, measuring and displaying
CC gene expression in samples derived from the human heart via microarrays.
CC By measuring gene expression, the probes are useful for predicting,
CC diagnosing, grading, staging, monitoring and prognosis diseases of the
CC human heart and vascular system e.g. cardiovascular disease,
CC hypertension, cardiac arrhythmias and congenital heart disease.
CC Note: The sequence data for this patent did not form part of the printed
CC specification, but was obtained in electronic format directly from WIPO
CC at [ftp://wipo.int/pub/published_pct_sequences](http://wipo.int/pub/published_pct_sequences).
XX
SQ Sequence 50 AA;
SQ Sequence 50 AA;

Query Match 1.4%; Score 14; DB 22; Length 50;
Best local Similarity 100.0%; Pred. No. 2.6e-06;
Matches 14; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 955 LPADSCAIMEGED 968
Db 19 LPADSCAIMEGED 32

CC brain. They can be used to measure gene expression in brain cell samples,
CC which may enable the diagnosis and improved treatment of nervous system
CC diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia,
CC epilepsy and cancers. The present sequence is a protein encoded by one of
CC the probes of the invention.

CC Sequence 50 AA;
ID AAM73033
ID AAM73033 standard; Protein; 50 AA.
XX
AC AAM73033;
XX
DT 06-NOV-2001 (first entry)
XX
DE Human bone marrow expressed probe encoded protein SEQ ID NO: 33339.
XX
KW Human; bone marrow expressed exon; gene expression analysis; probe;
KW microarray; cancer; leukaemia; lymphoma; myeloma.
XX
OS Homo sapiens.
XX
PN WO200157276-A2.
XX
PD 09-AUG-2001.
XX
PP 30-JAN-2001; 2001WO-US00668.
XX
PR 04-FEB-2000; 2000US-0180312.
PR 26-MAY-2000; 2000US-0207456.
PR 30-JUN-2000; 2000US-0208408.
PR 03-AUG-2000; 2000US-0032366.
PR 21-SEP-2000; 2000US-0234687.
PR 27-SEP-2000; 2000US-0236359.
PR 04-OCT-2000; 2000GB-0024263.

XX
PA (MOLE-) MOLECULAR DYNAMICS INC.
XX
PT Penn SG, Hanzel DK, Chen W, Rank DR;
XX
DR WPI; 2001-488900/53.

XX
PT Human genome-derived single exon nucleic acid probes useful for
PT analyzing gene expression in human bone marrow -
XX
PS Example 4; SEQ ID NO: 33339; 650pp + Sequence Listing; English.
XX
CC The present invention provides a number of single exon nucleic acid
CC probes which are derived from genomic sequences expressed in the human
CC bone marrow. They can be used to measure gene expression in bone marrow
CC samples, which may enable the improved diagnosis and treatment of cancers
CC such as lymphoma, leukaemia and myeloma. The present sequence is a
CC protein encoded by one of the probes of the invention.

XX
SQ Sequence 50 AA;
SQ Sequence 50 AA;
Query Match 1.4%; Score 14; DB 22; Length 50;
Best local Similarity 100.0%; Pred. No. 2.6e-06;
Matches 14; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 955 LPADSCAIMEGED 968
Db 19 LPADSCAIMEGED 32

PT Single exon nucleic acid probes for analyzing gene expression in human
PT brains -
XX
PS Example 4; SEQ ID NO: 32502; 650pp + Sequence Listing; English.
XX
CC The present invention provides a number of single exon nucleic acid
CC probes which are derived from genomic sequences expressed in the human

PD	30-JAN-2001;	2001WO-US00665.
XX		RESULT 30
XX	AAU21145	AAU21145 standard; Protein; 71 AA.
PR	ID	AAU21345;
PR	XX	AAU21345;
PR	AC	AAU21345;
PR	XX	AAU21345;
PR	XX	18-DEC-2001 (first entry)
PR	XX	DE
PR	XX	Human novel foetal antigen, SEQ ID NO 1589.
PA	XX	KW Human; foetal tissue antigen; antiinflammatory; neuroprotective; immunomodulator; cardiovascular; cytotoxic; nephrotoxic; cardiovascular; autoimmune disease; rheumatoid arthritis;
PI	XX	KW hyperproliferative disorder; breast neoplasm; cancer; cardiovascular disorder; cardiac arrest; cerebrovascular disorder; cerebral ischaemia; angiogenesis; nervous system disorder; wound healing; epithelial cell proliferation; food additive.
Penn SG, Hanzel DK, Chen W, Rank DR;	XX	
XX	PS	
WPI; 2002-114183/15.	XX	
XX	PT	
Spatially-addressable set of single exon nucleic acid probes, used to measure gene expression in human lung samples -	XX	
XX	PT	
XX	PS	
Claim 27; SEQ ID No 32542; 634pp; English.	XX	
CC	CC	The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived from human lung comprising single exon nucleic acid probes having one of 12614 nucleic acid sequences mentioned in the specification, or their complements or the 12387 open reading frames derived from the 12614 probes. Also included are a microarray comprising the novel set of probes; the novel set of probes which hybridise at high stringency to a nucleic acid expressed in the human lung; measuring gene expression in a sample derived from human lung, comprising (a) contacting the array with a collection of detectably labeled nucleic acids derived from human lung mRNA, and (b) measuring the label detectably bound to each probe of the array; identifying exons in a eukaryotic genome comprising
CC	CC	(a) algorithmically predicting at least one exon from genomic sequences of the eukaryote; and (b) detecting specific hybridisation of detectably labeled nucleic acids from a eukaryote lung mRNA, to a single exon probe, having a fragment identical to the predicted exon, the probe is included in the above mentioned microarray; assigning exons to a single gene, comprising (a) identifying exons from genomic sequence by the method above and (b) measuring the expression of each of the exons in several tissues and/or cell types using hybridisation of the exon to a single exon microarrays having a probe with the exon, where a common pattern of expression of the exons in the tissues and/or cell types indicates that the exons should be assigned to a single gene; a peptide comprising one of 10/11 sequences, mentioned in the specification, or encoded by the probes/open reading frames (ORF). The probes are used for gene expression analysis, and for identifying exons in a gene, particularly using human lung derived mRNA and for the study of lung diseases such as asthma, lung cancer, chronic obstructive pulmonary disease (COPD), interstitial lung disease (ILD), familial idiopathic pulmonary fibrosis, neurofibromatosis, tuberous sclerosis, Gaucher's disease, Niemann-Pick disease, Hermansky-Pudlak syndrome, sarcoidosis, pulmonary haemangioderosis, pulmonary histiocytosis, lymphangioleiomyomatosis, pulmonary alveolar proteinosis, Kartagener syndrome, fibrocytic pulmonary dysplasia, primary ciliary dyskinesia, pulmonary hypertension and hyaline membrane disease. The present sequence is a peptide/protein encoded by a single exon probe of the invention.
CC	CC	Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp://ipo.int/pub/published_pct_sequences.
XX	SQ	Sequence 50 AA;
Oy	955	Query Match 1.4%; Score 14; DB 23; Length 50; Best local Similarity 100.0%; Pred. 2.6e-06; Matches 14; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

PR 06-SEP-2000; 2000US-0230438. PR 17-NOV-2000; 2000US-0249244.
 PR 08-SEP-2000; 2000US-0231242. PR 17-NOV-2000; 2000US-0249254.
 PR 08-SEP-2000; 2000US-0231244. PR 17-NOV-2000; 2000US-0249255.
 PR 08-SEP-2000; 2000US-0231413. PR 17-NOV-2000; 2000US-0249257.
 PR 08-SEP-2000; 2000US-0231414. PR 17-NOV-2000; 2000US-0249259.
 PR 08-SEP-2000; 2000US-0232080. PR 17-NOV-2000; 2000US-0249300.
 PR 08-SEP-2000; 2000US-0232081. PR 01-DEC-2000; 2000US-0251050.
 PR 12-SEP-2000; 2000US-0232401. PR 01-DEC-2000; 2000US-0251051.
 PR 14-SEP-2000; 2000US-0232397. PR 05-DEC-2000; 2000US-0251050.
 PR 14-SEP-2000; 2000US-0233063. PR 05-DEC-2000; 2000US-0251988.
 PR 14-SEP-2000; 2000US-0233065. PR 05-DEC-2000; 2000US-0256719.
 PR 14-SEP-2000; 2000US-0233239. PR 06-DEC-2000; 2000US-0251479.
 PR 14-SEP-2000; 2000US-02332401. PR 08-DEC-2000; 2000US-0251866.
 PR 14-SEP-2000; 2000US-0233064. PR 08-DEC-2000; 2000US-0251869.
 PR 14-SEP-2000; 2000US-0233065. PR 08-DEC-2000; 2000US-0251989.
 PR 14-SEP-2000; 2000US-0233239. PR 08-DEC-2000; 2000US-0251990.
 PR 21-SEP-2000; 2000US-0234274. PR 11-DEC-2000; 2000US-0254097.
 PR 25-SEP-2000; 2000US-0234997. PR 05-JAN-2001; 2001US-0259678.
 PR 25-SEP-2000; 2000US-0234998. XX
 PR 26-SEP-2000; 2000US-0235484. (HUMA-) HUMAN GENOME SCI INC.
 PR 27-SEP-2000; 2000US-0235834. XX
 PR 27-SEP-2000; 2000US-0235836. PI Rosen CA, Barash SC, Ruben SM;
 PR 29-SEP-2000; 2000US-0236367. XX
 PR 29-SEP-2000; 2000US-0236368. DR WPI; 2001-488782/53.
 PR 29-SEP-2000; 2000US-0236370. XX
 PR 02-OCT-2000; 2000US-0237037. PT N-PSDB; AA534165.
 PR 02-OCT-2000; 2000US-0237038. XX
 PR 02-OCT-2000; 2000US-0237040. PT
 PR 13-OCT-2000; 2000US-0239935. PT
 PR 13-OCT-2000; 2000US-0239937. PT
 PR 20-OCT-2000; 2000US-0240960. PT
 PR 20-OCT-2000; 2000US-0241221. PT
 PR 20-OCT-2000; 2000US-0241785. PT
 PR 20-OCT-2000; 2000US-0241787. PT
 PR 20-OCT-2000; 2000US-0241808. PT
 PR 20-OCT-2000; 2000US-0241809. PT
 PR 01-NOV-2000; 2000US-0241826. PT
 PR 08-NOV-2000; 2000US-0244617. PT
 PR 08-NOV-2000; 2000US-0244647. PT
 PR 08-NOV-2000; 2000US-02446475. PT
 PR 08-NOV-2000; 2000US-02446476. PT
 PR 08-NOV-2000; 2000US-0246477. PT
 PR 08-NOV-2000; 2000US-0246478. PT
 PR 08-NOV-2000; 2000US-0246523. PT
 PR 08-NOV-2000; 2000US-0246524. PT
 PR 08-NOV-2000; 2000US-0246525. PT
 PR 08-NOV-2000; 2000US-0246526. PT
 PR 08-NOV-2000; 2000US-0246527. PT
 PR 08-NOV-2000; 2000US-0246528. PT
 PR 08-NOV-2000; 2000US-0246532. PT
 PR 08-NOV-2000; 2000US-0246609. PT
 PR 08-NOV-2000; 2000US-0246610. PT
 PR 08-NOV-2000; 2000US-0246611. PT
 PR 08-NOV-2000; 2000US-0246613. PT
 PR 17-NOV-2000; 2000US-0249207. PT
 PR 17-NOV-2000; 2000US-0249208. PT
 PR 17-NOV-2000; 2000US-0249209. PT
 PR 17-NOV-2000; 2000US-0249215. PT
 PR 17-NOV-2000; 2000US-0249216. PT
 PR 17-NOV-2000; 2000US-0249217. PT
 PR 17-NOV-2000; 2000US-0249218. PT

Query Match 1.4%; Score 14; DB 22; Length 71;
 Best Local Similarity 100.0%; Pred. No. 3.6e-06; Mismatches 0; Indels 0; Gaps 0;
 Matches 14; Conservative 0; MisMatches 0; Indels 0; Gaps 0;

QY 955 1PAA6SCAIMGED 968
 DB 16 1PAA6SCAIMGED 29

RESULT 31
 ID AAB70285
 XX AAB70285 standard; peptide; 78 AA.

AC AAB70285; PF 20-MAR-2000; 2000WO-EP02478.
 XX DT XX
 10-MAY-2001 (first entry) PR 26-MAR-1999; 99GB-0007113.
 XX PR 25-SEP-1999; 99GB-002285B.
 DE XX PA (SMRK) SMITHKLINE BEECHAM BIOLOGICALS.
 XX KW PT Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 XX XX DR WPI; 2000-664923/64.
 OS XX PN Novel CASB619 polyPeptides useful for diagnosing, and as vaccines for
 XX PD prophylactic and therapeutic treatment of, cancers, particularly
 XX ovarian and colon carcinoma, and autoimmune diseases -
 XX PT Example 7; Page 62; 68pp; English.
 XX PR XX The present sequence comprises an epitope derived from the human CASB619
 CC protein sequence. This protein is thought to be specifically or
 CC over-expressed in tumour cells, and so can be used as a target for
 CC antigen-specific immune responses which can cause destruction of the
 CC tumour cell. In addition, the protein and gene can be used in cancer
 CC diagnosis, in the treatment of autoimmune diseases and in vaccines
 CC against cancer and autoimmune disease. The present sequence can be used
 CC as an immunogen.
 XX SQ Sequence 10 AA;
 XX DR Query Match 1.0%; Score 10; DB 21; Length 10;
 XX CC Best Local Similarity 100.0%; Pred. No. 0; 0; 0; 0; 0;
 CC PT Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 CC XX Db 1 AISSCVAGI 10
 PS Sequence 78 AA;
 XX SQ RESULT 33
 XX AAB27115 ID AAB27115 Standard; Protein; 10 AA.
 XX AC AAB27115;
 XX DT 12-FEB-2001 (first entry)
 XX DE Human CASB619 protein epitope SEQ ID NO: 41.
 XX KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine;
 XX Homo sapiens.
 XX OS XX Homo sapiens.
 XX PN WO200058460-A2.
 XX PD 05-OCT-2000.
 XX PP 20-MAR-2000; 2000WO-EP02478.
 XX PR 26-MAR-1999; 99GB-0007113.
 XX PR 25-SEP-1999; 99GB-002285B.
 XX PA (SMRK) SMITHKLINE BEECHAM BIOLOGICALS.
 XX PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 XX DR WPI; 2000-664923/64.
 XX PT Novel CASB619 polyPeptides useful for diagnosing, and as vaccines for
 XX PT prophylactic and therapeutic treatment of, cancers, particularly
 XX PT ovarian and colon carcinoma, and autoimmune diseases -
 XX XX Example 7; Page 62; 68pp; English.

The present sequence comprises an epitope derived from the human CASB61 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

```

Query Match          1 0%; Score 10; DB 21; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.01; Mismatches 0;
Matches 10; Conservative 0; Indels 0; Gaps 0;
Y 852 SAACAPLCSV 861
| 1 SAACAPLCSV 10
XX

```

ID NO: 43.
disease; immunogen; vaccine;

RESULT 34
 AAB27116
 AAB27116 standard; protein; 10 AA.
 AAB27116;
 12-FEB-2001 (first entry)

Human CASB619 protein epitope SEQ ID NO: 42.
 Human; CASB619; cancer; autoimmune disease; immunogen; vaccine; epitope.
 Homo sapiens.
 W6200058460-A2.
 05-OCT-2000.
 20-MAR-2000; 2000MO-EP02478.
 26-MAR-1999; 99GB-0007113.
 25-SEP-1999; 99GB-0022858.
 (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 WPT; 2000-664923/64.
 Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases -
 Example 7; Page 62; 68pp; English.
 The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

Sequence 10 AA;
 Query Match 1.0%; Score 10; DB 21; Length 10;
 Best Local Similarity 100.0%; Pred. No. 0.01; Mismatches 0; Indels 0; Gaps 0;
 Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

RESULT 35
 AAB27118 standard; Protein; 10 AA.
 AAB27118;
 12-FEB-2001 (first entry)

Human CASB619 protein epitope SEQ ID NO: 44.
 Human; CASB619; cancer; autoimmune disease; immunogen; vaccine; epitope.
 Homo sapiens.

Query Match 1.0%; Score 10; DB 21; Length 10;
 Best Local Similarity 100.0%; Pred. No. 0.01; Mismatches 0; Indels 0; Gaps 0;
 Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

RESULT 36
 AAB27118 standard; Protein; 10 AA.
 AAB27118;
 12-FEB-2001 (first entry)

Human CASB619 protein epitope SEQ ID NO: 44.
 Human; CASB619; cancer; autoimmune disease; immunogen; vaccine; epitope.

PT prophylactic and therapeutic treatment of, cancers, particularly
 PN ovarian and colon carcinoma, and autoimmune diseases -
 XX
 PD Example 7; Page 62; 68pp; English.

XX
 PF 20-MAR-2000; 2000WO-EP02478.
 XX
 PR 26-MAR-1999; 99GB-0007113.
 PR 25-SEP-1999; 99GB-0022858.
 XX
 PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 XX
 PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 XX
 DR WPI; 2000-664923/64.

PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for
 PT prophylactic and therapeutic treatment of, cancers, particularly
 PT ovarian and colon carcinoma, and autoimmune diseases -
 XX
 PS Example 7; Page 62; 68pp; English.

XX
 CC The present sequence comprises an epitope derived from the human CASB619
 CC protein sequence. This protein is thought to be specifically or
 CC over-expressed in tumour cells, and so can be used as a target for
 CC antigen-specific immune responses which can cause destruction of the
 CC tumour cell. In addition, the protein and gene can be used in cancer
 CC diagnosis, in the treatment of autoimmune diseases and in vaccines
 CC against cancer and autoimmune disease. The present sequence can be used
 CC as an immunogen.

XX
 CC Sequence 10 AA;

Query Match 1.0%; Score 10; DB 21; Length 10;
 Best local Similarity 100.0%; Pred. No. 0.01; Mismatches 0; Indels 0; Gaps 0;
 Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 761 SLADRLIGHT 770
 Db 1 SLADRLIGHT 10

RESULT 37

AAB27119
 ID AAB27119 standard; Protein; 10 AA.
 XX
 AC AAB27119;
 XX
 DT 12-FEB-2001 (first entry)
 XX
 DE Human CASB619 protein epitope SEQ ID NO: 46.
 XX
 KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine;
 KW epitope.
 XX
 OS Homo sapiens.
 XX
 PN WO20058460-A2.
 XX
 PD 05-OCT-2000.
 XX
 PR 20-MAR-2000; 2000WO-EP02478.
 XX
 PR 26-MAR-1999; 99GB-0007113.
 PR 25-SEP-1999; 99GB-0022858.
 XX
 PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 XX
 DR WPI; 2000-664923/64.

PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for
 PT prophylactic and therapeutic treatment of cancers, particularly
 PT ovarian and colon carcinoma, and autoimmune diseases -
 XX
 PS Example 7; Page 63; 68pp; English.

XX
 CC The present sequence comprises an epitope derived from the human CASB619
 CC protein sequence. This protein is thought to be specifically or
 CC over-expressed in tumour cells, and so can be used as a target for
 CC antigen-specific immune responses which can cause destruction of the
 CC tumour cell. In addition, the protein and gene can be used in cancer
 CC diagnosis, in the treatment of autoimmune diseases and in vaccines
 CC against cancer and autoimmune disease. The present sequence can be used
 CC as an immunogen.

XX
 CC Sequence 10 AA;

Query Match 1.0%; Score 10; DB 21; Length 10;

PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 XX
 DR WPI; 2000-664923/64.
 XX
 PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases -
 XX
 PS Example 7; Page 63; 68pp; English.
 XX
 CC The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.
 XX
 SQ Sequence 10 AA;

Query Match 1.0%; Score 10; DB 21; Length 10;
 Best Local Similarity 100.0%; Pred. No. 0.01; 0; Mismatches 0; Indels 0; Gaps 0;
 Matches 10; Conservative 0; OS Homo sapiens.
 Qy 778 ITSPAEFLHL 787
 Db 1 ITSPAEFLHL 10

RESULT 42

AAB27124
 ID AAB27124 standard; Protein; 10 AA.
 XX
 AC AAB27124;
 XX
 DT 12-FEB-2001 (first entry)
 XX
 DE Human CASB619 protein epitope SEQ ID NO: 51.
 XX
 Human; CASB619; cancer; autoimmune disease; immunogen; vaccine; epitope.
 KW XX
 PA Homo sapiens.
 OS XX
 PN WO20005460-A2.
 XX
 PD 05-OCT-2000.
 XX
 PF 20-MAR-2000; 2000WO-EP02478.
 XX
 PR 26-MAR-1999; 99GB-0007113.
 PR 25-SEP-1999; 99GB-0022858.
 XX
 PA (SMK) SMITHKLINE BEECHAM BIOLOGICALS.
 PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 XX
 DR WPI; 2000-664923/64.
 XX
 PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases -
 XX
 PS Example 7; Page 63; 68pp; English.
 XX
 CC The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.
 XX
 SQ Sequence 10 AA;

Query Match 1.0%; Score 10; DB 21; Length 10;
 Best Local Similarity 100.0%; Pred. No. 0.01; 0; Mismatches 0; Indels 0; Gaps 0;
 Matches 10; Conservative 0; OS Homo sapiens.
 Qy 428 TLPTNMETIV 437
 Db 1 TLPTNMETIV 10

RESULT 44

AAB27125
 ID AAB27125 standard; Protein; 10 AA.
 XX

The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

CC over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

SQ Sequence 10 AA;

Query Match 1.0%; Score 10; DB 21; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.01; Mismatches 0; Indels 0; Gaps 0;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 644 GTKNKHKSL 653
Db 1 GTKNKHKSL 10

RESULT 47

AAB27130
ID AAB27130 standard; Protein; 10 AA.

AC AAB27130;
XX DT 12-FEB-2001 (first entry)

XX DE Human CASB619 protein epitope SEQ ID NO: 57.

XX KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine; epitope.

XX OS Homo sapiens.

XX PN WO200058460-A2.

XX PD 05-OCT-2000.

XX PP 20-MAR-2000; 2000WO-EP02478.

XX PR 26-MAR-1999; 99GB-0007113.

XX PR 25-SEP-1999; 99GB-0022858.

XX PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.

XX PT Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;

XX DR WPI; 2000-664923/64.

XX PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases -

XX PS Example 7, Page 64; 68pp; English.

XX CC The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for

CC antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

XX SQ Sequence 10 AA;

Query Match 1.0%; Score 10; DB 21; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.01; Mismatches 0; Indels 0; Gaps 0;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 260 LVNIAITGV 269
Db 1 LVNIAITGV 10

RESULT 49

AAB27132
ID AAB27132 standard; Protein; 10 AA.

AC AAB27132;
XX DT 12-FEB-2001 (first entry)

XX DE Human CASB619 protein epitope SEQ ID NO: 58.

XX KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine; epitope.

XX OS Homo sapiens.

XX PN WO200058460-A2.

Query Match 1.0%; Score 10; DB 21; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.01; Mismatches 0; Indels 0; Gaps 0;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 465 SDNPFMLT 474
Db 1 SDNPFMLT 10

Sequence „

RESULT 52

QY 795 VIFFYRNDV 804
 ||||| OS
 ||||| Homo sapiens.
 ||||| XX
 ||||| PN WO200058460-A2.
 ||||| XX
 ||||| PD 05-OCT-2000.
 ||||| XX
 ||||| PR 20-MAR-2000; 2000WO-EP02478.
 ||||| XX
 ||||| PR 26-MAR-1999; 99GB-0007113.
 ||||| PR 25-SEP-1999; 99GB-0022858.
 ||||| XX
 ||||| PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 ||||| XX
 ||||| PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 ||||| XX
 ||||| DR WPI; 2000-664923/44.
 ||||| XX
 ||||| PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for
 ||||| PT prophylactic and therapeutic treatment of, cancers, particularly
 ||||| PT ovarian and colon carcinoma, and autoimmune diseases -
 ||||| XX
 ||||| PS Example 7; Page 65; 68pp; English.
 ||||| XX
 ||||| CC The present sequence comprises an epitope derived from the human CASB619
 ||||| CC protein sequence. This protein is thought to be specifically or
 ||||| CC over-expressed in tumour cells, and so can be used as a target for
 ||||| CC antigen-specific immune responses which can cause destruction of the
 ||||| CC tumour cell. In addition, the protein and gene can be used in cancer
 ||||| CC diagnosis, in the treatment of autoimmune diseases and in vaccines
 ||||| CC against cancer and autoimmune disease. The present sequence can be used
 ||||| CC as an immunogen.
 ||||| XX
 ||||| SQ Sequence 10 AA;
 ||||| XX
 ||||| Query Match 1.0%; Score 10; DB 21; Length 10;
 ||||| Best Local Similarity 100.0%; Pred. No. 0.01; 0.01%;
 ||||| Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 ||||| XX
 ||||| QY 712 SVCTNDNTDL 721
 ||||| XX
 ||||| DB 1 SVCTNDNTDL 10
 ||||| XX
 ||||| AC AAB27137;
 ||||| XX
 ||||| DT 12-FEB-2001 (first entry)
 ||||| XX
 ||||| DE Human CASB619 protein epitope SEQ ID NO: 63.
 ||||| XX
 ||||| KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine;
 ||||| KW epitope.
 ||||| XX
 ||||| Homo sapiens.
 ||||| XX
 ||||| PN WO200058460-A2.
 ||||| XX
 ||||| PD 05-OCT-2000.
 ||||| XX
 ||||| PR 20-MAR-2000; 2000WO-EP02478.
 ||||| XX
 ||||| PR 26-MAR-1999; 99GB-0007113.
 ||||| PR 25-SEP-1999; 99GB-0022858.
 ||||| XX
 ||||| PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 ||||| XX
 ||||| PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 ||||| XX
 ||||| KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine;
 ||||| KW epitope.

DR WPI; 2000-664923/64.

XX

PT

The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

Example 7; Page 65; 68pp; English.

Query Match 1.0%; Score 10; DB 21; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.01; Mismatches 0; Indels 0; Gaps 0;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 432 NMETTVLISGI 441
Db 1 NMETTVLISGI 10

RESULT 56
ID AAB27139 standard; Protein; 10 AA.
XX
AC AAB27139;
XX
DT 12-FEB-2001 (first entry)
XX
DE Human CASB619 protein epitope SEQ ID NO: 65.
XX
KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine; epitope.
XX
OS Homo sapiens.
XX
PN WO200058460-A2.
XX
PD 05-OCT-2000.
XX
PF 20-MAR-2000; 2000WO-EP02478.
XX
PR 26-MAR-1999; 99GB-0007113.
XX
ER 25-SEP-1999; 99GB-0022858.
XX
PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
XX
PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
XX
DR WPI; 2000-664923/64.
XX
PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases

Example 7; Page 65; 68pp; English.

PT The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

XX
SQ Sequence 10 AA;

Query Match 1.0%; Score 10; DB 21; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.01; Mismatches 0; Indels 0; Gaps 0;
Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 263 NIAITGVAYT 272
Db 1 NIAITGVAYT 10

RESULT 57
ID AAB27140
XX
AC AAB27140;

DT 12-FEB-2001 (first entry)
 XX Human CASB619 protein epitope SEQ ID NO: 66.
 DE Human; CASB619; cancer; autoimmune disease; immunogen; vaccine;
 KW epitope.
 XX Homo sapiens.
 OS WO20058460-A2.
 XX 05-OCT-2000.
 PD XX 20-MAR-2000; 2000WO-EP02478.
 PF XX 26-MAR-1999; 99GB-0007113.
 PR XX 25-SEP-1999; 99GB-0022858.
 (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 XX DR WPI; 2000-664923/64.
 DR Novel CASB619 polypeptides useful for diagnosing, and as vaccines for
 prophylactic and therapeutic treatment of, cancers, particularly
 ovarian and colon carcinoma, and autoimmune diseases
 XX PS Example 7; Page 66; 68pp; English.
 XX The present sequence comprises an epitope derived from the human CASB619
 protein sequence. This protein is thought to be specifically or
 over-expressed in tumour cells, and so can be used as a target for
 antigen-specific immune responses which can cause destruction of the
 tumour cell. In addition, the protein and gene can be used in cancer
 diagnosis, in the treatment of autoimmune diseases and in vaccines
 against cancer and autoimmune disease. The present sequence can be used
 as an immunogen.
 XX SQ Sequence 10 AA;
 DR Query Match 1.0%; Score 10; DB 21; Length 10;
 DR Best Local Similarity 100.0%; Pred. No. 0.01; Mismatches 0; Indels 0; Gaps 0;
 DR Matches 10; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 DR Qy 846 FHFILWESAA 855
 DR Db 1 FHFILWESAA 10
 DR ID AAB26181
 DR AC AAB26181;
 DR XX 12-FEB-2001 (first entry)
 DR Human CASB619 protein epitope SEQ ID NO: 4.
 DR XX Human; CASB619; cancer; autoimmune disease; immunogen; vaccine;
 DR KW epitope.
 DR XX Homo sapiens.
 DR OS WO20058460-A2.
 DR XX 05-OCT-2000.
 DR PF XX 20-MAR-2000; 2000WO-EP02478.
 DR PR XX 26-MAR-1999; 99GB-0007113.
 DR PR XX 25-SEP-1999; 99GB-0022858.
 DR PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 DR PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 DR XX DR WPI; 2000-664923/64.
 DR Novel CASB619 polypeptides useful for diagnosing, and as vaccines for
 prophylactic and therapeutic treatment of, cancers, particularly
 ovarian and colon carcinoma, and autoimmune diseases
 XX PS Example 7; Page 57; 68pp; English.
 XX The present sequence comprises an epitope derived from the human CASB619
 protein sequence. This protein is thought to be specifically or
 over-expressed in tumour cells, and so can be used as a target for
 antigen-specific immune responses which can cause destruction of the

CC tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

SQ Sequence 9 AA:

Query Match 0.9%; Score 9; DB 21; Length 9;
Best Local Similarity 100.0%; Pred No. 7.8e+05;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 24 RLWRLLWA 32
Db 1 RLWRLLWA 9

RESULT 60

AAB26182 AAB26182 standard; Protein; 9 AA.

AC AAB26182;

XX DT 12-FEB-2001 (first entry)

XX DE Human CASB619 protein epitope SEQ ID NO: 7.

XX KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine; epitope.

XX OS Homo sapiens.

XX PN WO200058460-A2.

XX PD 05-OCT-2000.

XX PF 20-MAR-2000; 2000WO-EP02478.

XX PR 26-MAR-1999; 99GB-0007113.

XX PR 25-SEP-1999; 99GB-0022858.

XX PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.

XX PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;

XX DR 2000-664923/64.

XX PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases -

XX PS Example 7; Page 57; 68pp; English.

CC The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically over-expressed in tumour cells, and so can be used as a target for CC antigen-specific immune responses which can cause destruction of the CC tumour cell. In addition, the protein and gene can be used in cancer CC diagnosis, in the treatment of autoimmune diseases and in vaccines CC against cancer and autoimmune disease. The present sequence can be used CC as an immunogen.

XX SQ Sequence 9 AA;

Query Match 0.9%; Score 9; DB 21; Length 9;
Best Local Similarity 100.0%; Pred No. 7.8e+05;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 893 SIEPEQRTVI 901
Db 1 SIEPEQRTVI 9

RESULT 62

AAB26184 AAB26184 standard; Protein; 9 AA.

AC AAB26184;

XX DT 12-FEB-2001 (first entry)

XX DE Human CASB619 protein epitope SEQ ID NO: 8.

XX KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine; epitope.

XX OS Homo sapiens.

XX PN WO200058460-A2.

XX PD 05-OCT-2000.

Query Match 0.9%; Score 9; DB 21; Length 9;
Best Local Similarity 100.0%; Pred No. 7.8e+05;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 761 SLADRLIGV 769
Db 1 SLADRLIGV 9

RESULT 61

PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases -

Example 7; Page 58; 68pp; English.

The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

SQ Sequence 9 AA;

Query Match 0.9%; Score 9; DB 21; Length 9;
Best Local Similarity 100.0%; Pred. No. 7.8e+05;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 903 KTIIDFWLK 9
ID AAB26192 standard; Protein; 9 AA.

Db 1 KTIIDFWLK 9

RESULT 69
AAB26191
ID AAB26191 standard; Protein; 9 AA.

AC AAB26191;
XX

DT 12-FEB-2001 (first entry)

XX Human CASB619 protein epitope SEQ ID NO: 15.

DB 1 KLEVKYKSL 944

OS Homo sapiens.

XX PN WO20058460-A2.

PD 05-OCT-2000.

XX PF 20-MAR-2000; 2000WO-EP02478.

XX PR 26-MAR-1999; 99GB-0007113.

XX PR 25-SEP-1999; 99GB-0022858.

XX PA (SMITH) SMITHKLINE BEECHAM BIOLOGICALS.

PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;

XX DR WPI; 2000-664923/64.

XX PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases -

XX PS Example 7; Page 58; 68pp; English.

XX The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or

over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the

tumour cell. In addition, the protein and gene can be used in cancer

diagnosis, in the treatment of autoimmune diseases and in vaccines

against cancer and autoimmune disease. The present sequence can be used as an immunogen.

XX SQ Sequence 9 AA;

Query Match 0.9%; Score 9; DB 21; Length 9;
Best Local Similarity 100.0%; Pred. No. 7.8e+05;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 860 SVADYHAIV 868
ID AAB26193 standard; Protein; 9 AA.

Db 1 SVADYHAIV 9

RESULT 70
AAB26193
ID AAB26193 standard; Protein; 9 AA.

AC AAB26193;
XX

DT 12-FEB-2001 (first entry)

XX SQ Sequence 9 AA;

The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

DE Human CASB619 protein epitope SEQ ID NO: 17.
 XX
 KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine;
 KW epitope.
 OS Homo sapiens.
 XX
 PN WO200058460-A2.
 PD 05-OCT-2000.
 XX
 PF 20-MAR-2000; 2000WO-EP02478.
 XX
 PR 26-MAR-1999; 99GB-0007113.
 PR 25-SEP-1999; 99GB-0022858.
 XX
 PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 XX
 Bruck CEM, Casbatt J, Coche T, Vinals De Bassols YC;
 XX
 PT
 XX
 DR
 XX
 PS
 XX
 PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for
 prophylactic and therapeutic treatment of, cancers, particularly
 ovarian and colon carcinoma, and autoimmune diseases -
 XX
 PS Example 7, Page 59; 68pp; English.
 XX
 CC The present sequence comprises an epitope derived from the human CASB619
 protein sequence. This protein is thought to be specifically or
 over-expressed in tumour cells, and so can be used as a target for
 antigen-specific immune responses which can cause destruction of the
 tumour cell. In addition, the protein and gene can be used in cancer
 diagnosis, in the treatment of autoimmune diseases and in vaccines
 against cancer and autoimmune disease. The present sequence can be used
 CC as an immunogen.
 XX
 SQ Sequence 9 AA;
 CC The present sequence comprises an epitope derived from the human CASB619
 protein sequence. This protein is thought to be specifically or
 over-expressed in tumour cells, and so can be used as a target for
 antigen-specific immune responses which can cause destruction of the
 tumour cell. In addition, the protein and gene can be used in cancer
 diagnosis, in the treatment of autoimmune diseases and in vaccines
 against cancer and autoimmune disease. The present sequence can be used
 CC as an immunogen.
 XX
 SQ Sequence 9 AA;
 CC The present sequence comprises an epitope derived from the human CASB619
 protein sequence. This protein is thought to be specifically or
 over-expressed in tumour cells, and so can be used as a target for
 antigen-specific immune responses which can cause destruction of the
 tumour cell. In addition, the protein and gene can be used in cancer
 diagnosis, in the treatment of autoimmune diseases and in vaccines
 against cancer and autoimmune disease. The present sequence can be used
 CC as an immunogen.
 XX
 RESULT 71
 AAB26194
 ID AAB26194 standard; Protein: 9 AA.
 XX
 AC AAB26194;
 XX
 DT 12-FEB-2001 (first entry)
 XX
 DE Human CASB619 protein epitope SEQ ID NO: 19.
 XX
 KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine;
 KW epitope.
 XX
 OS Homo sapiens.
 XX
 PN WO200058460-A2.
 XX
 PD 05-OCT-2000.
 XX
 PF 20-MAR-2000; 2000WO-EP02478.
 XX
 PR 26-MAR-1999; 99GB-0007113.
 PR 25-SEP-1999; 99GB-0022858.
 XX
 PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 XX
 Bruck CEM, Casbatt J, Coche T, Vinals De Bassols YC;
 XX
 PT
 XX
 DR
 XX
 PS Example 7, Page 59; 68pp; English.
 XX
 CC The present sequence comprises an epitope derived from the human CASB619
 protein sequence. This protein is thought to be specifically or
 over-expressed in tumour cells, and so can be used as a target for
 antigen-specific immune responses which can cause destruction of the
 tumour cell. In addition, the protein and gene can be used in cancer
 diagnosis, in the treatment of autoimmune diseases and in vaccines

CC against cancer and autoimmune disease. The present sequence can be used
CC as an immunogen.

XX

Sequence 9 AA;

SQ

Query Match Best Local Similarity 0.9%; Score 9; DB 21; Length 9;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 503 TLCSYNC1 511
Db 1 TLCSYNC1 9

RESULT 73

AAB26196

standard; Protein; 9 AA.

XX

AC AAB26197;

XX

DT 12-FEB-2001 (first entry)

XX

DB Human CASB619 protein epitope SEQ ID NO: 21.

XX

KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine;

KW epitope.

XX

OS Homo sapiens.

XX

PN WO20058460-A2.

XX

PD 05-OCT-2000.

XX

PF 20-MAR-2000; 2000WO-EP02478.

XX

PR 26-MAR-1999; 99GB-0007113.

XX

PR 25-SEP-1999; 99GB-0022858.

XX

PA (SMK) SMITHKLINE BEECHAM BIOLOGICALS.

XX

PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;

XX

DR WPI; 2000-664923/64.

XX

PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for ovarian and colon carcinoma, and autoimmune diseases

XX

PS Example 7, Page 59, 68pp; English.

XX

CC The present sequence comprises an epitope derived from the human CASB619

CC protein sequence. This protein is thought to be specifically or

CC over-expressed in tumour cells, and so can be used as a target for

CC antigen-specific immune responses which can cause destruction of the

CC tumour cell. In addition, the protein and gene can be used in cancer

CC diagnosis, in the treatment of autoimmune diseases and in vaccine

CC against cancer and autoimmune disease. The present sequence can be used

CC as an immunogen.

XX

SQ Sequence 9 AA;

XX

Qy 169 NTDPBP1 177

Db 1 NTDPBP1 9

RESULT 74

AAB26197

ID AAB26197 standard; Protein; 9 AA.

XX

The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccine against cancer and autoimmune disease. The present sequence can be used as an immunogen.

Example 7, Page 59, 68pp; English.

Novel CASB619 polypeptides useful for diagnosing, and as vaccines for ovarian and colon carcinoma, and autoimmune diseases

Example 7, Page 59, 68pp; English.

The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccine against cancer and autoimmune disease. The present sequence can be used as an immunogen.

Sequence 9 AA;

Query Match Best Local Similarity 0.9%; Score 9; DB 21; Length 9;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 81 SLPDPVKG 89
Db 1 SLPDPVKG 9

RESULT 75

AAB26199
ID AAB26199 standard; Protein; 9 AA.

AC AAB26199;
XX

DT 12-FEB-2001 (first entry)

XX

DB Human CASB619 protein epitope SEQ ID NO: 23.

XX

KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine;

KW epitope.

XX

OS Homo sapiens.

XX

PN WO20058460-A2.

XX

PR 05-OCT-2000.

XX

PT 20-MAR-2000; 2000WO-EP02478.

Tue Apr 22 16:18:05 2003

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XX 99GB-0007113.
 PR 99GB-0022858.
 XX
 PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 XX
 PT Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 PI
 XX
 DR 2000-664923/64.
 XX
 Novel CASB619 polypeptides useful for diagnosis, and as vaccines for
 PT prophylactic and therapeutic treatment of, cancers, particularly
 PT ovarian and colon carcinoma, and autoimmune diseases -
 XX
 PS Example 7; Page 59; 68pp; English.

XX The present sequence comprises an epitope derived from the human CASB619
 PS protein sequence. This protein is thought to be specifically or
 CC over-expressed in tumour cells, and so can be used as a target for
 CC the over-expressed protein sequence. This protein is thought to be specifically or
 CC over-expressed in tumour cells, and so can be used as a target for
 CC antigen-specific immune responses which can cause destruction of the
 CC tumour cell. In addition, the protein and gene can be used in vaccines
 CC against cancer and autoimmune disease. The present sequence can be used
 CC as an immunogen.

XX Query Match Similarity 100.0%; Pred. No. 7.8e+05;
 DR Best Local Similarity 0%; Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Query Match Similarity 100.0%; Pred. No. 7.8e+05;
 DR Best Local Similarity 0%; Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

XX Sequence 9 AA;
 PS 0.9%; Score 9; DB 21; Length 9;
 DR 0.9%; Score 9; DB 21; Length 9;
 CC Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 CC Mismatches 0; Indels 0; Gaps 0;
 CC Matches 9; Conservative 9;
 XX
 OY 867 IVSCVAGI 875
 DB 1 IVSCVAGI 9

CC The present sequence comprises an epitope derived from the human CASB619

RESULT 78

AAB26202

ID

XX

AC

AAB26202;

XX

DT

12-FEB-2001 (first entry)

XX

DE

Human

KW

XX

KW

Human; CASB619; protein epitope SEQ ID NO: 26.

KW

epitope.

XX

OS

Homo sapiens.

XX

PN

W0200058460-A2.

XX

PD

05-OCT-2000.

XX

PR

20-MAR-2000; 2000WO-EP02478.

XX

PR

26-MAR-1999; 99GB-0007113.

XX

PR

25-SEP-1999; 99GB-0022858.

XX

PA

(SMIK) SMITHKLINE BEECHAM BIOLOGICALS.

XX

PI

Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;

XX

DR

WPI; 2000-664923/64.

XX

PT

Novel CASB619 polypeptides useful for diagnosing, and as vaccines for

PT

prophylactic and therapeutic treatment of, cancers, particularly

XX

PS

ovarian and colon carcinoma, and autoimmune diseases

XX

CC

The present sequence comprises

CC

an epitope derived from the human CASB619

CC

protein sequence. This protein is thought to be specifically or

CC

over-expressed in tumour cells, and so can be used as a target for

CC

antigen-specific immune responses which can cause destruction of the

CC

tumour cell. In addition, the protein and gene can be used in the

CC

diagnosis, in the treatment of autoimmune diseases and in vaccines

CC

against cancer and autoimmune disease. The present sequence can be used

CC

as an immunogen.

CC

as an immunogen, and autoimmune disease. The present sequence can be used

CC

as an immunogen.

KW Human; **CASB619**; cancer; autoimmune disease; immunogen; vaccine;
KW epitope.
OS *Homo sapiens*.
XX
PN WO200058460-A2.
XX
PD 05-OCT-2000.
XX
PP 20-MAR-2000; 2000WO-EP02478.
XX
PR 26-MAR-1999; 99GB-0007113.
XX
PR 25-SEP-1999; 99GB-0022858.
PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
XX
PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
XX
DR WPI; 2000-664923/64.
PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases -
XX
PS Example 7; Page 61; 68pp; English.
CC The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.
XX
SQ Sequence 9 AA;
Query Match 0.9%; **Score** 9; **DB** 21; **Length** 9;
Best Local Similarity 100.0%; **Pred. No.** 7.8e+05;
Matches 9; **Conservative** 0; **Mismatches** 0; **Indels** 0; **Gaps** 0;
Db 1 GISDPEQRV 9
Qy 914 SAGCTTAT 922
Db 1 SAGCTTAT 9
RESULT 84
AC AAB27106;
ID AAB27106 standard; Protein; 9 AA.
XX
AC AAB27106;
ID AAB27106 standard; Protein; 9 AA.
XX
DT 12-FEB-2001 (first entry)
XX
DE Human CASB619 protein epitope SEQ ID NO: 33.
XX
KW Human; **CASB619**; cancer; autoimmune disease; immunogen; vaccine;
XX
OS *Homo sapiens*.
XX
PN WO200058460-A2.
XX
PD 05-OCT-2000.
XX
PP 20-MAR-2000; 2000WO-EP02478.
XX
PR 26-MAR-1999; 99GB-0007113.
XX
PR 25-SEP-1999; 99GB-0022858.
PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
XX
PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
XX
PR 26-MAR-1999; 99GB-0007113.
XX
PR 25-SEP-1999; 99GB-0022858.
PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
XX
PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
XX
PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for prophylactic and therapeutic treatment of, cancers, particularly ovarian and colon carcinoma, and autoimmune diseases -
XX
PS Example 7; Page 61; 68pp; English.
CC The present sequence comprises an epitope derived from the human CASB619 protein sequence. This protein is thought to be specifically or over-expressed in tumour cells, and so can be used as a target for antigen-specific immune responses which can cause destruction of the tumour cell. In addition, the protein and gene can be used in cancer diagnosis, in the treatment of autoimmune diseases and in vaccines against cancer and autoimmune disease. The present sequence can be used as an immunogen.

XX Sequence . 9 AA;
 SQ 0.9%; Score 9; DB 21; Length 9;
 Query Match Best Local Similarity 100.0%; Pred. No. 7.8e+05; Mismatches 0;
 Matches 9; Conservative 0; Indels 0; Gaps 0;

QY 824 KTVFGSLL 832
 Db 1 KTVFGSLL 9

RESULT 86
 AAB27108 DT 12-FEB-2001 (first entry)
 ID AAB27108 XX Human CASS619 protein epitope SEQ ID NO: 35.
 XX DE Human; CASS619; cancer; autoimmune disease; immunogen; vaccine;
 XX KW epitope.
 XX OS Homo sapiens.
 XX PN WO20058460-A2.

XX PD 05-OCT-2000.
 XX PF 20-MAR-2000; 2000WO-BP02478.
 XX PR 26-MAR-1999; 99GB-0007113.
 XX PR 25-SEP-1999; 99GB-0022858.
 XX PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 XX PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 XX DR WPI; 2000-664923/64.
 XX PT Novel CASS619 polypeptides useful for diagnosing, and as vaccines for
 PT prophylactic and therapeutic treatment of, cancers, particularly
 PT ovarian and colon carcinoma, and autoimmune diseases -
 XX PS Example 7; Page 61; 68pp; English.

XX CC The present sequence comprises an epitope derived from the human CASS619
 CC protein sequence. This protein is thought to be specifically or
 CC over-expressed in tumour cells, and so can be used as a target for
 CC antigen-specific immune responses, which can cause destruction of the
 CC tumour cell. In addition, the protein and gene can be used in cancer
 CC diagnosis, in the treatment of autoimmune diseases and in vaccines
 CC against cancer and autoimmune disease. The present sequence can be used
 CC as an immunogen.

XX SQ Sequence . 9 AA;
 CC Query Match 0.9%; Score 9; DB 21; Length 9;
 PT Best Local Similarity 100.0%; Pred. No. 7.8e+05; Mismatches 0;
 PT Matches 9; Conservative 0; Indels 0; Gaps 0;

QY 681 TLAGGPST 689
 Db 1 TLAGGPST 9

RESULT 88
 AAB27110 DT 12-FEB-2001 (first entry)
 ID AAB27110 XX Human CASS619 protein epitope SEQ ID NO: 36.
 XX DE Human; CASS619; cancer; autoimmune disease; immunogen; vaccine;
 XX KW epitope.
 XX OS Homo sapiens.
 XX PN WO20058460-A2.

XX PD 05-OCT-2000.
 XX PF 20-MAR-2000; 2000WO-BP02478.
 XX PR 26-MAR-1999; 99GB-0007113.
 XX AC AAB27109

RESULT 87
 AAB27109 DT 12-FEB-2001 (first entry)
 ID AAB27109 XX Human CASS619 protein epitope SEQ ID NO: 35.
 XX KW epitope.
 XX OS Homo sapiens.
 XX PN WO20058460-A2.

XX PD 05-OCT-2000.
 XX PF 20-MAR-2000; 2000WO-BP02478.
 XX PR 26-MAR-1999; 99GB-0007113.
 XX AC AAB27109;

PR 25-SEP-1999; 99GB-0022858.
 XX
 PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 XX
 PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 XX
 DR WPI; 2000-664923/64.
 XX
 PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for
 ovarian and colon carcinoma, and autoimmune diseases -
 XX
 PS Example 7; Page 61; 68pp; English.
 The present sequence comprises an epitope derived from the human CASB619
 protein sequence. This protein is thought to be specifically or
 over-expressed in tumour cells, and so can be used as a target for
 antigen-specific immune responses which can cause destruction of the
 tumour cell. In addition, the protein and gene can be used in cancer
 diagnosis, in the treatment of autoimmune diseases and in vaccines
 against cancer and autoimmune disease. The present sequence can be used
 as an immunogen.
 XX
 SQ Sequence 9 AA;

Query Match 0.9%; Score 9; DB 21; Length 9;
 Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 539 YIEEENNTT 547
 Db 1 YIEEENNTT 9

RESULT 89

AAB27111
 ID AAB27111 standard; Protein; 9 AA.
 XX
 AC AAB27111;
 XX
 DT 12-FEB-2001 (first entry)
 XX
 DE Human CASB619 protein epitope SEQ ID NO: 38.
 XX
 KW Human; CASB619; cancer; autoimmune disease; immunogen; vaccine;
 XX
 OS Homo sapiens.
 XX
 PN WO200058460-A2.
 XX
 PD 05-OCT-2000.
 XX
 PP 20-MAR-2000; 2000WO-EP02478.
 XX
 PR 26-MAR-1999; 99GB-0007113.
 XX
 PR 25-SEP-1999; 99GB-0022858.
 XX
 PA (SMIK) SMITHKLINE BEECHAM BIOLOGICALS.
 XX
 PI Bruck CEM, Cassart J, Coche T, Vinals De Bassols YC;
 XX
 DR WPI; 2000-664923/64.
 XX
 PT Novel CASB619 polypeptides useful for diagnosing, and as vaccines for
 ovarian and colon carcinoma, and autoimmune diseases -
 XX
 PS Example 7; Page 61; 68pp; English.
 The present sequence comprises an epitope derived from the human CASB619
 protein sequence. This protein is thought to be specifically or
 over-expressed in tumour cells, and so can be used as a target for
 antigen-specific immune responses which can cause destruction of the
 tumour cell. In addition, the protein and gene can be used in cancer
 diagnosis, in the treatment of autoimmune diseases and in vaccines
 against cancer and autoimmune disease. The present sequence can be used
 as an immunogen.
 XX
 SQ Sequence 9 AA;

Query Match 0.9%; Score 9; DB 21; Length 9;
 Best Local Similarity 100.0%; Pred. No. 7.8e+05;
 Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 38 VTOGTGPEL 46
 Db 1 VTOGTGPEL 9

The present sequence comprises an epitope derived from the human CASB619
 protein sequence. This protein is thought to be specifically or
 over-expressed in tumour cells, and so can be used as a target for

PA (MOLE-) MOLECULAR DYNAMICS INC.
PT Penn SG, Hanzel DK, Chen W, Rank DR
XX DR WPI; 2001-488899/53.
PT Single exon nucleic acid probes for analyzing gene expression in human
XX hearts -
PS Claim 15; SEQ ID No 25666; 530pp; English.
CC The present invention relates to single exon nucleic acid probes for
CC measuring human gene expression in a sample derived from human heart (see
CC AB21535 AB41305). The present sequence is a protein encoded by one suc-
CC gene expression in samples derived from the human heart via microarrays.
CC BY measuring gene expression, the probes are useful for predicting, measuring and displaying
CC diagnosing, grading, staging, monitoring and prognosis diseases of the
CC human heart and vascular system e.g. cardiovascular disease,
CC hypertension, cardiac arrhythmias and congenital heart disease,
CC Note: The sequence, cardiac arrhythmias and congenital heart disease,
CC specification, but was obtained in electronic format directly from WIPO
XX at http://wipo.int/pub/published_pct_sequences.
Sequence 32 AA;
Query Match 0.9%; Score 9; DB 22; Length 32;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0
Qy 964 MEGEDVDD 972
Db 1 MEGEDVDD 9
RESULT 94
ID AAM59554
XX AAM59554 standard; Protein; 32 AA. -
AC AAM59554;
XX
DT 05-NOV-2001 (first entry)
DE Human brain expressed single exon probe encoded protein SEQ ID NO: 31659.
KW Human; brain expressed exon; gene expression analysis; probe;
KW microarray; Alzheimer's disease; multiple sclerosis; schizophrenia;
KW epilepsy; cancer.
OS Homo sapiens.
XX
PN WO20015275-A2.
XX
PD 09-AUG-2001.
XX
PP 30-JAN-2001; 2001WO-US00667.
PR 04-FEB-2000; 2000US-0180312.
PR 26-MAY-2000; 2000US-0207456.
PR 30-JUN-2000; 2000US-0008408.
PR 03-AUG-2000; 2000US-0532356.
PR 21-SEP-2000; 2000US-0234657.
PR 27-SEP-2000; 2000US-0236339.
PR 04-OCT-2000; 2000GB-0024263.
PA (MOLE-) MOLECULAR DYNAMICS INC.
PI Penn SG, Hanzel DK, Chen W, Rank DR
XX DR WPI; 2001-483446/52.
PT Single exon nucleic acid probes for analyzing gene expression in human
XX brains -

KW Human; bone marrow expressed exon; gene expression analysis; probe;
 KW microarray; cancer; leukaemia; lymphoma; myeloma.
 XX
 OS Homo sapiens.
 XX
 PN WO200157276-A2.
 XX
 PD 09-AUG-2001.
 XX
 PP 30-JAN-2001; 2001WO-US00668.
 PR 04-FEB-2000; 2000US-0180312.
 PR 26-MAY-2000; 2000US-0207456.
 PR 30-JUN-2000; 2000US-0608408.
 PR 03-AUG-2000; 2000US-0632466.
 PR 21-SEP-2000; 2000US-0234687.
 PR 27-SEP-2000; 2000US-0236359.
 PR 04-OCT-2000; 2000GB-0024263.
 PA (MOL-3) MOLECULAR DYNAMICS INC.
 XX
 PI Penn SG, Hanzel DK, Chen W, Rank DR;
 XX
 DR WPI; 2001-48890/53.
 PT Human genome-derived single exon nucleic acid probes useful for
 PT analyzing gene expression in human bone marrow -
 XX
 PS Example 4; SEQ ID NO: 32429; 658pp + Sequence Listing; English.
 The present invention provides a number of single exon nucleic acid
 probes which are derived from genomic sequences expressed in the human
 bone marrow. They can be used to measure gene expression in bone marrow
 samples, which may enable the improved diagnosis and treatment of cancer
 such as lymphoma, leukaemia and myeloma. The present sequence is a
 protein encoded by one of the probes of the invention.
 Sequence 32 AA;
 KK

us-10-046-433-40.oligo.rag

RESULT 100
 AAM78689
 ID AAM78689 standard; Protein; 748 AA..
 AC
 XX
 DT 06-NOV-2001 (first entry)
 DE Human protein SEQ ID NO 1351.

KW Human; cytokine; cell proliferation; cell differentiation; gene therapy;
 KW vaccine; peptide therapy; stem cell growth factor; haematopoiesis;
 KW tissue growth factor; immunomodulatory; cancer; leukaemia;
 KW nervous system disorder; arthritis; inflammation.
 OS Homo sapiens.
 XX
 PN WO200157190-A2.
 XX
 PD 09-AUG-2001.
 XX
 PF 05-FEB-2001; 2001WO-US04098.
 XX
 PR 03-FEB-2000; 2000US-0496914.
 PR 27-APR-2000; 2000US-0560875.
 PR 20-JUN-2000; 2000US-0598075.
 PR 01-SEP-2000; 2000US-0620325.
 PR 15-SEP-2000; 2000US-0663561.
 PR 20-OCT-2000; 2000US-0693325.
 PR 30-NOV-2000; 2000US-0728422.
 XX
 PA (HYSE-) HYSQ INC.
 XX
 PI Tang YT, Liu C, Drmanac RT, Asundi V, Zhou P, Xu C, Cao Y, Ma Y;
 PI Zhao QA, Wang D, Wang J, Zhang J, Ren F, Chen R, Wang ZW;
 PI Xue AJ, Yang Y, Wejhrman T, Goodrich R;
 DR WPI: 2001-176283/51.
 DR N-PSDB; AAK51822.

XX
 PT Nucleic acids encoding polypeptides with cytokine-like activities,
 PT useful in diagnosis and gene therapy
 XX
 PS Claim 20; Page 3596-3597; 6221pp; English.
 XX
 CC The invention relates to polynucleotides (AAK51456-AAK53455) and the
 CC encoded polypeptides (AAM8323-AAM80321) that exhibit activity relating to
 CC cytokine, cell proliferation or cell differentiation or which may induce
 CC production of other cytokines in other cell populations. The
 CC polynucleotides and polypeptides are useful in gene therapy, vaccines or
 CC peptide therapy. The polypeptides have various cytokine-like activities,
 CC e.g. stem cell growth factor activity, haematopoiesis regulating
 CC activity, tissue growth factor activity, immunomodulatory activity and
 CC activin/inhibin activity and may be useful in the diagnosis and/or
 CC treatment of cancer, leukaemia, nervous system disorders, arthritis and
 CC inflammation.
 Note: Records for SEQ ID NO 2110 (AAK52581), 2111 (AAK52582) and 3666
 CC (AAM8020) are omitted as the relevant pages from the sequence listing
 CC were missing at the time of publication.
 XX
 SQ Sequence 748 AA;

Query Match 0.9%; Score 9; DB 22; Length 748;

Best Local Similarity 100.0%; Pred. No. 5.6;
 Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 964 MEGEDVEDD 972
 Db 8 MEGEDVEDD 16

Search completed: April 22, 2003, 15:33:47
 Job time : 88 secs

